

350	
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AGGATGTTGA CATCTGATCG CGACACCGAA CTAATAGGAC CATAGGTGTC AATTCCACTA	18360
ACATAGATAT TGAAAGACTG ACTCTTAGAC GTCTTAGGAG CTTCTACTTT TTTAGTGAAT	18420
CCCTTAGTAT AAATCTTTT TATCTCGAT GCGTAGTCTG GATACTCTGA CTCGATGATG	18480
TTTTCAAAGA CACTATTTAG GACAATGGCC TTAGTCTCCC CTGCAATCAA ACTCTTGAA	18540
GCTGCCAAGT AAGACGAACT CTGGTTGACC GTCAAATCGG TATTCTGACT TGACTTGATA	18600
TCAGCTAGTA ATTTCTGAAT ATTTTCATTA TTAGTCCCAG TCGGTGCTGT CACACTCGTC	18660
AGTTGCGTAA CATTTCGAT CTCACTATCT GCTAAAACAG CGACACTGAT TGAATATTCT	18720
GAGTAATTAG AAGTCGCATT TAAACGATTG GTCAGTCCAA CAAACTGCTG TACTGCAAAG	18780
AGCGACACAG AGCTGACAAG GATAGAGAAC ACCAACAGAA AAATAGTAAA CTTTCAGCT	18840
TTTTTATAGA TAATCAAGAG TAGCCCTACC AAGGCAACTA GTAGGACTAA CGCAGTTACC	18900
ACTAGATTAAGATATCTAAA AGCAAGGATA TTGTACTTAA AGATTAAGAA CAATAAAAAAA	18960
CAAACATAACA ATAATAAAAT AGTCAGCAAA ACTATATTAA CACTTCGTT CACTTTCTGT	19020
GAACGTGATT TTTTAAACG TCTACTCATG ATTAATACCT ATACATTGAA CATTATAAGGA	19080
TTATATCACT TTTTACGGT AATGTCTACA CCTTTATTTT TACTATCTGC ATCTTTAAGT	19140
ATCTTAGTAG ACTTCCCGCG AAACAAAAAT ATAGTAAAT GAAATAAGAA CAGAACAAAT	19200
CGTTCAAGGAC ACTCAAATCG ATTCTCAACA ATGTTTTAGA AGCAGAGGTG	19250

(2) INFORMATION FOR SEQ ID NO: 36:

- (i) SEQUENCE CHARACTERISTICS:
- (A) LENGTH: 21706 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 36:

AAAGTTGAAA GACTGCTAGC TGTTTTGAT ACCAATCGTT TCCAACTACA GAGCAAACAG	60
TATACAAAAGT TTGTTTTGG ATGTAAGCTT CTTGATGGAC AATTCCAAGA AAATCAAGAA	120
ATTGCTGACC TTCAATTTT TGCCATTGAC CAACTGCCGA ACTTATCTGA AAAACGCATT	180
ACCAAGGAGC AAATAGAGCT TCTTTGGCAG GTTTATCAAG GTCATAGGGG GCAATATCTT	240

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GACTAAGAAG ATGATTATCG TATTCTAAA TCCATTTTA ACAACTAGCA TGGTATAATA	300
ATATGCAGGA AAATTTGAA TTATGAGGAA GACTAGATGA ATTATGGGA TATTTCCTT	360
ACGACTCAGG CAACCGAGCC GCCCAAATTT GACCTTTTT GGTATGTTAG CCTATTTACG	420
CTCTTAGCCT TAACCTTTA TACAGCCAT CGCTATCGTG AAAAGAAGGT TTACCAACGA	480
TTTTCCAAA TCTTCAGAC TGTTCAAGTA ATCCTCTTT ATGGTTGGTA CTGGGTCAAT	540
CATATGCCAC TGTCAGAAAG CCTACCCATT TACCATTGCC GTATGGCTAT GTTTGTGGTA	600
CTCTTGCTTC CTGGTCAATC CAAATATAAA CAATACTTG CATTATTGGG AACATTTGGG	660
ACATTAGCAG CCTTGTGTTA TCCAGTGCCA GATGCTTACC CTTTCCACA TATCACCAT	720
CTATCCTTTA TCTTGGTCA TTTAGCACTC TTGGGAACT CTCTAGTTA TCTATTGAGA	780
CA GTATAATG CGCGATTGCT GGATGTGAAG GGAATTTTC TCATGACCTT TGCCCTAAAT	840
GCCTTGATTT TTGTTGCAAA TTGGTACA GGTGGCGATT ACGGATTTT GACAAAACCG	900
CCATTGGTTG GGGATCACGG TCTAGTAGCT AATTATTTAC TTGTTCAAT TGTGCTGGTA	960
GCTACTATCA GTTGTACTAA GAAAATCTTA GAATTCTTT TAGCTCAAGA AGCAGAAAAA	1020
ATGATTGCAA AGGAAGCTTA ACACAGAGCT TTCTTTTTG CTCTTAGAGA GTTTTACAA	1080
GCAGCTTATA AAATAAGAAT TTCTGAATAG ACAAACTCAA AAAATGGCTG GGAAATTTAG	1140
GAAAAAAAGCA AGCACCGATTA AATTTTTGT GTTATAATAT TTTGTGAATA GCTATGCCA	1200
TGTTTAGCTA TCGAAATAATA CGAAGTGCAGA AACTTGGAAAG ATAGAGAGGA AGCGATGTAA	1260
TGGCTAGAGA AGGCTTTTT ACAGGTCTAG ATATTGGAAC AAGCTCTGTC AAGGTGCTTG	1320
TGGCCGAGCA GAGAAATGGT GAATTAAATG TAATTGGCGT GAGTAATGCC AAAAGTAAAG	1380
GTGTAAAGGA TCGAAATTATT GTTGATATTG ATGCAGCAGC AACTGCTATC AAGTCAGCCA	1440
TTTCCCAAGC GGAAGAAAAG GCAGGCATTT CGATTAAATC AGTGAATGTC GGCTTGCTG	1500
GTAATCTTT GCAGGTAGAA CCAACTCAGG GGATGATTCC AGTAACATCT GATACTAAGG	1560
AAATTACGGA TCAAGATGTT GAAAATGTTG TCAAATCAGC TTTGACAAAG AGTATGACAC	1620
CTGACCGTGA AGTCATTACC TTATTCCTG AAGAATTAT TGTGGATGGT TTCCAAGGG	1680
TTCGTGACCC ACGTGGCATG ATGGGGGTTG GCCTTGAAAT GCGTGGTTG CTTTATACAG	1740
GACCTCGTAC TATCTGCAC AATTGCGTA AGACGGTTGA GCGTGCAGGT GTTCAGGTTG	1800
AAAATGTTAT CATTTCACCA CTAGCAATGG TTCAGTCTGT TTTGACAAAG GGGGAACGTG	1860
AATTGGTGC TACAGTGATT GATATGGGGG CAGGTCAAAC GACTGTCGCT ACAATCCGTA	1920
ATCAAGAACT CCAGTCACA CATATTCTCC AAGAAGGTGG AGATTATGTA ACTAAAGATA	1980

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TCTCCAAGGT TTTGAAAACC TCTCGCAAAT TAGCGGAAGG CTTGAAACTG AATTACGGGG	2040
AAGCCTATCC GCCTCTTGCA AGCAAAGAAA CCTTCCAAGT AGAGGTTATT GGAGAAGTAG	2100
AAGCAGTCGA AGTCACGGAA GCCTACTTG T CAGAAATTAT TTCTGCACGA ATCAAGCACA	2160
TCCTTGAACA AATCAAGCAA GAATTAGATA GAAGGCCTCT ATTGGACCTC CCTGGTGGTA	2220
TTGTCTTAAT CGGTGGGAAT GCCATTTAC CAGGTATGGT TGAGCTTGCT CAGGAAGTCT	2280
TTGGCGTCG TGTCAAGCTT TATGTTCCAA ATCAAGTTGG TATCCGTAAT CCAGCCTTTG	2340
CGCATGTGAT TAGTTTATCA GAATTGCGG GTCAATTAAAC AGAAGTTAAT CTTTTGGCTC	2400
AGGGAGCGAT AAAAGGTGAG AATGACTTAA GTCATCAGCC AATTAGTTTT GGTGGGATGC	2460
TGCAAAAAAC AGCTCAGTTT GTACAATCAA CGCCTGTTCA ACCAGCTCCT GCTCCAGAAC	2520
TAGAGCCGGT GGCGCCTACA GAACCAATGG CGGATTTCAC ACAAGCTTCA CAAAATAAAC	2580
CGAAATTAGC AGATCGTTTC CGTGGATTGA TCGGAAGCAT GTTTGACGAA TAAAGAGGAA	2640
AAATAAAATTA TGACATTTTC ATTTGATACA GCTGCTGCTC AAGGGGCAGT GATTAAAGTA	2700
ATTGGTGTG TGGGAGGTGG TGCAATGCC ATCAACCGTA TGGTCGACGA AGGTGTTACA	2760
GGCGTAGAAT TTATCGCAGC AAACACAGAT GTACAAGCAT TGAGTAGTAC AAAAGCTGAG	2820
ACTGTTATTC AGTTGGGACC TAAATTGACT CGTGGTTTG GTGCAGGAGG TCAACCTGAG	2880
GTTGGTCGTA AAGCCGCTGA AGAAAGCGAA GAAACACTGA CGGAAGCTAT TAGGGTGCC	2940
GATATGGTCT TCATCACTGC TGGTATGGGA GGAGGCTCTG GAACTGGAGC TGCTCCTGTT	3000
ATTGCTCGTA TCGCCAAAGA TTTAGGTGCG CTTACAGTTG GTGTTGTAAC ACgtCCCTTT	3060
GGTTTTGAAG GAAGTAAGCG TGGACAATTG GCTGTAGAAG GAATCAATCA ACTTCGTGAG	3120
CATGTAGACA CTCTATTGAT TATCTCAAAC AACAAATTGC TTGAAATTGT TGATAAGAAA	3180
ACACCGCTT TGGAGGCTCT TAGCGAAGCG GATAACGTTC TTCGTCAAGG TGTCAAGGG	3240
ATTACCGATT TGATTACCAA TCCAGGATTG ATTAACCTTG ACTTTGCCA TGTGAAAAGC	3300
GTAATGGCAA ACAAAAGGGAA TGCTCTTATG GGTATTGGTA TCGGTAGTGG AGAAGAACCT	3360
GTGGTAGAAG CGGCACGTTAA GGCAATCTAT TCACCACTTC TTGAAACAAC TATTGACGGT	3420
GCTGAGGATG TTATCGTCAA CGTTACTGGT GGTCTTGACT TAACCTTGAT TGAGGCAGAA	3480
GAGGCTTCAC AAATTGTGAA CCAGGCAGCA GGTCAAGGAG TGAACATCTG GCTCGGTACT	3540
TCAATTGATG AAAGTATGCG TGATGAAATT CGTGTAAACAG TTGTTGCAAC GGGTGTTCGT	3600
CAAGACCGCG TAGAAAAGGT TGTGGCTCCA CAAGCTAGAT CTGCTACTAA CTACCGTGAG	3660
ACAGTGAAAC CAGCTCATTC ACATGGCTTT GATCGTCATT TTGATATGGC AGAAACAGTT	3720
GAATTGCCAA AACAAAATCC ACGTCGTTTG GAACCAACTC AGGCATCTGC TTTTGGTGAT	3780

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TGGGATCTTC	GCCGTGAATC	GATTGTTCGT	ACAACAGATT	CAGTCGTTTC	TCCAGTCGAG	3840
CGCTTTGAAG	CCCCAATTC	ACAAGATGAA	GATGAATTGG	ATACACCTCC	ATTTTTCAAA	3900
AATCGTTAACG	TAATGAATG	TAAAAGAAAA	TACAGAACTT	GTTCGGTCTCT	AAGTTGCAGA	3960
GGCTAGTCTG	AGTGCTCATC	GAGAGAGTGG	TTCGGTCTCT	GTCATTGCAG	TTACCAAGTA	4020
TGTAGATGTA	CCGCACAGCGG	AAGCCTTGCT	TCGGCTAGGT	GTCCATCATA	TCGGTGAAAA	4080
TCGTGTAGAT	AAGTTCTGG	AAAAATATGA	AGCTTTAAAA	GATCGAGATG	TGACTTGGCA	4140
TTTGATTGGT	ACCTTGCAAA	GACGTAAGGT	AAAAGATGTC	ATTCAATACC	TTGATTATT	4200
CCATGCATTG	GACTCAGTAA	AGCTAGCAGG	GGAAATTCAA	AAAAGAAGTC	ACCGAGTCAT	4260
CAAGTGTTC	CTTCAGTAA	ATATTCCTAA	AGAAGAAAGC	AAACACGGTT	TTTCGAGAGA	4320
GGAAC TGCTG	GAATCTTGC	CAGAGTTAGC	CAGACTAGAT	AAGATTGAAT	ATGTTGGTT	4380
AATGACGATG	GCACCTTTG	AGGCTAGCAG	TGAGCAGTTG	AAAGAGATTT	TCAAGGCAG	4440
CCAAGATTTA	CAAAGAGAAA	TTCAAGAGAA	ACAAATTCCA	AAATAGCCTA	TGACCGAGTT	4500
AAGTATGGGA	ATGAGTCGTG	ATTATAAAGA	AGCGATTCAA	TTCGGTTCCA	CTTTGTTCG	4560
TATAGGTACA	TCATTTTTA	AGTAGGAGAG	AACCATGTCT	TTAAAGATA	GATTCGATAG	4620
ATTTATAGAT	TATTTTACGG	AGGATGAGGA	TTCAAGTCTC	CCTTATGAAA	AAAGAGATGA	4680
GCCTGTGTTT	ACTTCAGTAA	ATTCTTCACA	GGAACCGGCT	CTCCCAATGA	ATCACCTTC	4740
ACAGTCGGCT	GGCACAAAAG	AGAACAAATAT	CACCAGACTT	CATGCAAGAC	AAACAGGAATT	4800
GCCAAATCAG	AGTCAGCGTG	CAACGGATAA	GGTCATTATA	GATGTTCGTT	ATCCTAGAAA	4860
ATATGAGGAT	GCAACAGAAA	TTGTTGATTT	ATTGGCAGGA	AACGAAAGTA	TCTTGATTGA	4920
TTTTCAGTAT	ATGACAGAGG	TGCAAGCTCG	TCGTTGTTG	GACTATTTGG	ATGGAGCTTG	4980
TCATGTTTA	GCTGAAATT	TGAAAAGGT	AGCTTCTACC	ATGTATTTGT	TGACACCAGT	5040
GAACGTTATT	GTAATGTTG	AAGATATCCG	TTTACCAAGAT	GAAGATCAAC	AGGGTGAGTT	5100
CGGTTTGAT	ATGAAGCGAA	ATAGAGTACG	ATAATGATTT	TTTTAATTG	TATGATTTAT	5160
AATGCAGTGG	ATATTTACTC	CCTGATTTG	GTAGCCTTCG	CTGTCATGTC	TTGGTTTCCA	5220
GGTGCCTACG	AATCCAGTTT	AGTCGTTGG	ATTGCTAGCGT	TGGTGAAACC	AGTCCTTGCT	5280
CCCTTGCAAC	GCCTGCCTT	ACAGATAGCG	GGTCTTGATT	TATCTGTTG	GGTTGCGATT	5340
GTTTGGTTC	GATTTTTAGG	AGAAAACCTA	GTGCGTTTC	TGGCGATGAT	AGGATGAATA	5400
AAGGGATTTA	TCAGCATTTC	TCCATAGAAG	ATCGTCCATT	TCTTGACAAG	GGAATGGAAT	5460
GGATAAAAGAA	GGTAGAAGAT	AGCTATGTC	CTTTTTAAC	TCCTTTATC	AATCCTCATC	5520

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AGGAGAAAGCT ATAAAGATT TTGGCCAAAA CCTATGGTCT TGCTTGAGC AGTAGTGGGG	5580
AATTCGTCTC GAGTGAGTAT GTTCGAGTTT TATTATAACCC AGATTATTTCAACCGAGT	5640
TTTCAGATTT TGAAATATCT CTCCAGGAAA TTGTGTATTC CAATAAATTGAACATTTAA	5700
CGCATGCTAA GATTTAGGG ACAGTCATCA ATCAATTAGG GATTGAACGG AAACTTTTG	5760
GAGATATCCT AGTAGATGAA GAACGGGCAG AGATTATGAT TAATCAGCAG TTTCTTCTTC	5820
TCTTCAAGA TGGACTAAAG AAAATTGGTC STATACTGT TTCGCTGGAG GAACGTCCTT	5880
TCACCGAGAA AATAGATAAG CTAGAACAGT ATCGAGAACT GGATTTATCT GTGTCTAGTT	5940
TTCGATTAGA TGTCTTTA TCAAATGTT TGAAACTATC TAGGAATCAA GCAAACCAAGT	6000
TGATTGAAAA GAAACTTGTCAAGTAAATT ATCATGTGGT AGACAAATCA GATTACACTG	6060
TTCAAGTTGG AGACTTGATT AGTGTGAGAA AATTTGGTCG CTTGAGATTA CTTCAAGATA	6120
AGGGACAAAC GAAAAAGAG AAGAAAAAAA TAACCGTCCA GTTATTATTA AGTAAGTGAG	6180
GAATAGAATGCCAATTACAT CATTAGAAAT AAAGGACAAG ACTTTGGAA CTCGATTCAAG	6240
AGGTTTGAT CCAGAAGAAG TCGATGAATT TTTAGATATT GTGGTTCGTG ATTACGAAGA	6300
TCTTGTGCGT GCGAATCATG ATAAAAATTGCGTATTAAG AGTTTAAAG AGCGTTTGTCAAG	6360
TTACTTTGAT GAAATAAAAG ATTCAATTGAG CCAGTCTGTA TTGATTGCTC AGGATACAGC	6420
TGAGAGAGTG AAACAGGCAG CGCATGAACG TTCAAACAAAT ATCATTCAATC AAGCAGAGCA	6480
AGATGCGCAA CGCTTGTGG AAGAAGCTAA ATATAAGGCA AACGAGATTCTTCGTTCAAGC	6540
AACTGATAAT GCTAAGAAAG TCGCTGTGA AACAGAAGAA TTGAAGAACAGAGCCGTGT	6600
CTTCCACCAA CGTCTCAAAT CTACAATTGA GAGTCAGTTG GCTATTGTG AATCTTCAGA	6660
TTGGAAAGAT ATTCTCCGTC CAACAGCTAC TTATCTCAA ACCAGTGATG AAGCCTTTAA	6720
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TGATATGACA CGTCAGTTCT CTCAGCAGA AATGGCAGAA TTACAAGCTC GTATTGAGGT	6840
AGCCGATAAA GAATTGTCTG AATTTGAAGC TCAGATTAAA CAGGAAGTGG AAGCTCCAAC	6900
TCCTGTAGTG AGTCTCAAG TTGAAGAAGA GCCTCTGCTC ATCCAGTTGG CCCAATGTAT	6960
GAAGAACCG AAGTAGCTCC AATGCATCCG ATAGGTCCAA CACCAAGCTAC AGAAACTGTT	7020
GATTCAATAC CGGGATTGAGCAGCAGA GAATCTGTTA CAATTTTATA AGAAATATTCA	7080
TGAGAACAAAT ATCTTATCCT TATATTTCAGA GCGAGCAGGA GATGGTGTGA GTCTGTAAAT	7140
CCCTATTGAT AAGATTATCC TCTCAAAAC TCAAGTCTGA AGCTAGTAAG ATTTGACGTT	7200
TCCCACGTTA CGGGATAAGA GGGAGAAAGA CTAATCTTT TTCCGAATAA AGGTGGTACC	7260
ACGATTTTCG TCCTTTTGG AAGTCGTGGT TTTTAATTG TTATTATTTA TAAAGGAGAT	7320

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ACCATGAAAC TCAAAGACAC CCTTAATCTT GGGAAAATC AATTCCCAAT GCGTGCAGGC	7380
CTTCCTACCA AAGAGCCAGT TTGGCAAAAG GAATGGGAAG ATGCAAAACT TTATCAACGT	7440
CGTCAAGAAT TGAACCAAGG AAAACCTCAT TTCACCTTGC ATGATGGCCC TCCATACGCT	7500
AACGGAAATA TCCACGTTGG ACATGCTATG ACAAGATTT CAAAAGATAT CATTGTTCGT	7560
TCTAACTCTA TGTCAGGATT TTACGCACCA TTTATTCCCTG GTTGGGATAC TCATGGCTG	7620
CCAATCGAGC AAGTCTTGTC AAAACAAGGT GTCAAACGTA AAGAAATGGA CTTGGTTGAG	7680
TACTTGAAC TTTGCCGTGA GTACGCTCTT TCTCAAGTAG ATAACAACG TGAAGATTT	7740
AAACGTTTGG GTGTTCTGG TGACTGGAA AATCCATATG TGACCTTGAC TCCTGACTAT	7800
GAAGCAGCTC AAATTCGTGT ATTGGTGAG ATGGCTAATA AGGGTTATAT CTACCGTGGT	7860
GCTAAGCCAG TTTACTGGTC ATGGTCATCT GAGTCAGCAC TTGCTGAAGC AGAGATTGAA	7920
TACCATGACT TGGTTCAAC TTCCCTTAC TATGCCAACA AGGTAAAAGA TGGCAAAGGA	7980
GTTCTAGATA CAGATACTTA TATCGTTGTC TGGACAACGA CTCCATTAC CATCACAGCT	8040
TCTCGTGGTT TGACGGTTGG TGCGAGATATT GATTACGTTT TGGTTCAACC TGCTGGTGA	8100
GCTCGTAAGT TTGTCGTTGC TGCTGAATTA TTGACTAGCT TGTCTGAGAA ATTTGGCTGG	8160
GCTGATGTTCAAGT TGGTCAAC TACAGCCCCGT GGCCAAAGAAC TCAACCACAT CGTAACAGAA	8220
CACCCATGGG ATACAGCTGT AGAAGAGTTG GTAATTCTTG GTGACCACGT TACGACTGAC	8280
TCTCGTACAG GTATTGTCCA TACAGCCCCGT GGTTTGGTG AGGACGATTA CAATGTTGGT	8340
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GCTGGTCCTG AATTGAGG TCAATTCTAT GAAAAGGTAG TTCCAACGT TATTGAAAAA	8460
CTTGGTAACC TCCTTCTTGC CCAAGAAGAA ATCTCTCACT CATATCCATT TGACTGGCGT	8520
ACTAAGAAAC CAATCATCTG GCGTGCAGTT CCACAATGGT TTGCTCAGT TTCTAAATTG	8580
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AAACAAATCT TGTACAAGG TTTGCCCTT GATGGTAAAG GTGAGAAGAT GTCTAAATCT	9120
CTTGGAAATA CTATTGCTCC AAGCGATGTT GAAAACAAT TCGGTGCTGA AATCTTGCCT	9180
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CAAGTTCTG AAACCTACCG TAAGATTCGT AACACTCTTC GTTCTTGAT TGCCAATACA	9300
TCTGACTTTA ACCCAGCTCA AGATACAGTC GCTTACGATG AGCTTCGTTG AGTTGATAAG	9360
TACATGACGA TTCGCTTAA CCAGCTTGTCA AAGACCATTG GTGATGCCA TGCGAGACTT	9420
GAATTCTTGA CGATCTACAA GGCTTGGTG AACTTTATCA ACCTTGACTT GTCAGCCTTC	9480
TACCTGATT TTGCCAAAGA TGTGTTTAC ATTGAAGGTG CCAAATCACT GGAACGCCGT	9540
CAAATGCAGA CTGTCTTCTA TGACATTCTT GTCAAAATCA CCAAACCTTT GACACCAATC	9600
CTTCCTCACA CTGCGGAAGA AATCTGGTCA TATCTTGAGT TTGAAACAGA AGACTTCGTC	9660
CAATTGTCAG AATTACCAAGA AGTCAAACT TTTGCTAACCC AAGAAAGAAAT CTTGGATACA	9720
TGGGCAGCCT TCATGGACTT TCGTGGACAA GCACAAAAAG CCTTGGAAAGA AGCTCGTAAT	9780
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GCGGAAGCAG TCGCAGAAGG ATTTGAAGAG AAATAAGATT GAAAAGTCTA GGCAAAATTC	10140
AATTTGAGAA GAAAAGACAA CTAATTTTAT AGTCTATTAA ACGCATTGTA TCACGTTTT	10200
GAATACCTGA TATGATGCGT TTTTATTTA TTTTAAATTT TTGCGAGGTG TGACTTTTTA	10260
TACTCAACAA GAATCAAAGA GAAACTTAGC AAGCTAACAG TAGTAAGATA AAATAGGAAT	10320
TTGATATTAG GGATAAGATT GGTAAATAGT GTAATATTTT TACAACAATA AATTTATATA	10380
GTTATTTCTG GTTCTGAAA AGTATTATAT TTTATTTCAT ATTATACAAA TTTTTATTT	10440
ATAATATCG AACATACTTT TTTTAAAGC AAATATGATA CAATTTATT TGAAAAAAAT	10500
AAAAAAGGAG ATTTTATTAT AAAATAAAA AGACTTGCTT TAATTAAGTGG TATCGTCGGT	10560
CTTGTGGAG GAATTTTACT TCTTATTGGT CCTTTTGCT TGTGGGAAT AGCGGTAAAC	10620
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CTCTTGAATG CCTTGAAGAT TGCAAATCTT GTTCTTGGTA TCATTGCTAT TGTTTACTAT	10740
AAAGGAGATA AGCGTGTAGG TGCAGCTCCG TCTGTACTAA TGATTGTTTC TGGTGGAGTT	10800
AGTCTCATTC TATTCCGTTTC TTAGGATGGG TTGGGGGGAT TTTTGCTATT ATCGGAGGAT	10860

CTCTATTCCCT TTCAACATTG AAGAAATTCA AATCAGAAGA ATAAAAGGT A TTTTAGCATG	10920
AAAAGAACAA AAAAGTTTAT CGGTATAGGA GTAGCTCTAT TATCTCTTC TCTTCTAGTT	10980
GCATGTGGAA CATAAAAGTTC AAAGAATACT TCAACAAGTA ATGATGAGAA GACAGTAGCA	11040
ACATCCAATA GTTCAAAAGA ACAAATCACT TTCGATACAC CGGTTGTAAC AGACGATGCG	11100
ATTGAATCAA TACGCACTTA TGAGATTAT ATAGATCTT ATAAAAATAT TTTTGATGAT	11160
TATTTTACTA AAGCTGAGGA AGGTTCAAA GGCATAGCTA TGAAAATAA TGACTCGTT	11220
ACTAAACTAA AAGAGTCAAC TCAAAAATTA TTGATGCC AGAAAAAAAG GTTAAATAAT	11280
GAAGATAGAA TAGAAACAC CAAAACAAT GTGATTGCC AACATTGTCA AACAGTCCTT	11340
TCCTTTTG G TTTGACTAG CTCTTTGTG AAAAATTGTG TAAAATAGAA TAGATAAACG	11400
AGGGGAAACC TCGGAAATT TAAAGGAGAA TCCATCTAA GGTAATTG GTTTTGCTC	11460
GCCACGGTGA GTCTGAATGG AACAAAGCTA ACCTTTCAC TGTTGGCT GATGTTGATT	11520
TGTCTGAAAA AGGTACACAA CAAGCGATTG ACGCTGGTAA ATTGATCAA GAAGCTGGTA	11580
TGAAATTGCA CCAAGCTTAC ACTTCAGTAT TGAAACGTGC TATCAAACACA ACTAACTMGG	11640
CTCTTGAAAGC TTCTGACCAA TTGTGGGTTG CAGTTGAAAA ATCATGGCGC TTGAACGAAC	11700
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ATGCTGAAAA CTTGAAAGTG ACTTTGGAAC GTGCTCTTCC ATTCTGGAA GATAAAATCG	11940
CTCCAGCTCT TAAAGATGGT AAAAACGTAT TCGTAGGAGC TCACGGTAAC TCAATCCGTG	12000
CCCTTGAAA ACACATCAA GGTTGTCAG ATGACGAGAT CATGGACGTG GAAATCCCTA	12060
ACTTCCCACC ATTGGTATTC GAATTCGACG AAAAATTGAA CGTCGTTCT GAATACTACC	12120
TTGGAAAATA AAAAATTGTA AGTCTAGAAT TGATTTCTAG` GCTTTTTATG TTAGTATGGA	12180
AGTATGATAA GGAATAAAA ACAAGATTAT GTACTGGCCT ACAAGCAACC AGCTTCAACC	12240
ACTTACATGG GTTGGGAAGA AGAAGCTTA CCCATAGGCA ATGTTCTTT AGGAGCAAAA	12300
GTATTTGGCC TTATAGGGC TGAACGGATT CAATTAAATG AAAAAGTCT CTGGTCTGGA	12360
GGTCCACTTC CTGATAGTTC AGATTATCAG GGTGGAAATC TTCAGGATCA GTATGTTTT	12420
TTAGCTGAGA TTCGGCAGGC TTTGGAGAAG AGAGATTACA ATCTGGCTAA GGAAGTGGCT	12480
GAGCAGCACC TAATTGGGCC AAAAACGAGT CAATATGGGA CCTATCTGTC TTTTGGGGAT	12540
ATTACACATTG AGTTCAAGCCA GCAAGGTACG ACTTTGTCTC AGGTGACGGG CTATCAGAGA	12600

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CAGCTGAATA TTAGTAAGGC ACTTGGCAGC ACTTCTTATG TCTATAAGGG AACGGCATTT	12660
GAACGTAAAG CTTTGCGAG TTTTCAGAT GATCTCTGG TTCATGTTT TACTAAGGAA	12720
GGGTTGGAAA CTCTAGATTT TACTATAGAA CTATCCTTGA CCTGTGATT GGCTTCTGAT	12780
GGAAAGTATG AGCAGGAAAA ATCTGATTAC AAGGAGTGTA AGTTGGATAT TACTGATTCT	12840
CATATCTTGA TGAAGGGAAG AGTTAAGGAT AATGATCTGC GGTTTGCTAG TTATCTAGCT	12900
TGGGAAACGG ATGGAGATAT TAGAGTTGG TCAGATAGGG TTCAGATATC AGGAGCCAGT	12960
TATGCCAATC TCTTCTTGGC CGCTAACGACG GATTTGCCA AAAATCCTGC TAGCAATTAT	13020
CGCAAGAAC TAGATTTAGA GCACACAGGTG ATAGACTTGG TGGACACAGC TAAAGAAAAG	13080
GGCTATACCC AATTGAAATC AAGGCATATC GAGGACTACC AAGCCTTATT CCAGCGTGT	13140
CAATTGGATT TGGAAAGCTGA TGGTGCAGCA TCCACTACAG ATGATTGTT AAAAAATTAT	13200
AAGCCAAG AAGGGCAGGC TTTGGAGGAG CTGTTCTTCC AGTATGGACC GTATTTATTG	13260
ATTAGTTCGT CCAGAGACTG CCCAGATGCT CTACCAAGCTA ACCTACAGGG AGTCTGGAAT	13320
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TATTGCCAG CCTATGTTAC CAATCTCCTA GAGACGGTCT TTCCAGTCAT CAACTATGTA	13440
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CCTGGTTGGG ATTACTATTG GGTTGGTCA CCAGCTGCCA ATGCGTGGAT GATGCAAACC	13620
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ATGTTGAGGG AAACGGTTCG TTTTGGAAAT GCCTTTTAC ATAAGGATCA GCAGGCGCAG	13740
CGTTGGGTGT CTTCTCCGTC TTATTCCCCA GAACATGGGC CGATTCGAT TGGCAATACC	13800
TATGACCAAT CTCTGATTG GCAGTTATTT CATGATTTCATC TTCAGGCTGC TCAGGAATTG	13860
GGACTGGATG AGGACTTGTG GACTGAGGTT AAGGAGAACT GTGATTACT AAATCCTTTG	13920
CAAATCACTC AATCTGGTCG AATCAGGGAG TGGTATGAGG AGGAAGAGCA GTATTTCAA	13980
AATGAGAAAG TGGAGGCCA GCATCGGCAC GCTTCCCAC TACTGGGACT CTATCCTGGC	14040
AATCTTTA GCTACAAGGG ACAAGAGTAT ATTGAAGCGG CGCGTGCTAG CCTCAATGAT	14100
CGTGGAGATG GCGGCACAGG CTGGTCCAAG GCTAATAAGA TCAATCTCG GGCGCGTTG	14160
GGAGATGGCA ATCGAGGCCA TAAATTATTG GCAGAGCACT TAAAGACATC CACCTTGCAA	14220
AATCTTGGT GTAGCCATCC TCCTTTTCAG ATAGATGGTA ATTTTGGTGC TACTAGTGGC	14280
ATGGCAGAAA TGTTACTCCA GTCTCATGCA GCTTATCTGG TACCTCTAGC TGCCCTACCT	14340
GATGCTTGGT CAACAGGTTTC TGTTTCAGGC TTAATGGCAC GTGGACATTT TGAAGTGAGC	14400

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TTGCGAGTTT CTTATCCAGA TATTGAGAAG AGTGTGATTA AAATGAATCA AGAAAAAAATA	14520
AAAGCAGAAAT GCATGGGAA AGATTGTATT TCGGTGGCAA CAGCAGAAGG TGATCTGTT	14580
CAATTTTATT TTTAAGAAGA TGTTATAAGG CAGTAATTG AAACGCCTT TTAATAAGGA	14640
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TGTTTGAGG TTGCAGATGG AAGCTGACGT GGTTTGAAGA GAGATTTTCG AGGAGTATAA	14820
TTTGTGAT AGAGGGTGGG TCTGATGGCT TATATTGAGA TGAAACACTG TTACAAGCGT	14880
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GTAGCCAAA ATCCTAAAAT TCTCCTTTGT GATGAACCGA CTGGAGCCTT GGATTATCAG	15360
ACGGCCAAGC AGGTTTGAA AATTCTCCAA GACATGTCTC GTCAAAAGGG AGCGACGGTG	15420
ATCATCGTGA CTCATAATGG AGCTTTGGCG CCCATTGCTG ATCGCGTGTAT TCAAATGCAC	15480
GATGCCAGTG TCAAGGATGT GGTGCTCAAC CAGCATCCTC AGGATATTGA CAGTTGGAG	15540
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CCTCAAAGTA ACCAGTCCCA ACATGGAGGC GACAGCTAAT GCTTATTAA CAACTGCTCA	15720
AACCTGGAT TTGGCAGTCAGTCAACTA TGGCTGGAT CAAGCAGACC AAGAAGAACT	15780
AAAACAGACG GAGGGCGCAG AGTCGAGTT TGGCTATTG ACAGATGTGA CTATGGATAA	15840
TGGGCAGGAT GCCATTCCGGC TGTACTCCAA ACCAGAGCGA ATTTCACCT TTCAGCTAAG	15900
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AGACCATACT TATACCATTA CTGGTTTGT GGATTGGCT GAAATCCTCT CCCAGCGAGA	16080
TATGGGCTAC GCAGGAAGTG GAAGTGGGAC TCTGACAGCC TATGGGTGA TTTTACCTAG	16140

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TCTAGACAAG GGGCAAGAGA CCCTTGACAA GGCTCAGACT AATTTGCAGG AAGGCAAGCG	16380
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TCAAGTTCAAG AGAGAGCAGG CTAGTGCCTCA ACTTACCCAA GCCAAGCAGG ATTGGGCAA	16500
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GACCATGCCA GGTGGTCAGG GCTATCTTAT GTATAGCAAT GCTTCATCCA GTATTCGAGC	16680
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CATGACTCGC TTGTTAGACG AAGAGCGAAC TCATGCAGGG ATTGTTAAGG CCTTGGGTTA	16800
TCGTAGTAAG GATATTATCG CCAAGTTCT CCTTTATGGA CTAGTAGCTG GGACTGTCGG	16860
AACGGCTCTA GGTAGTATAC TTGGTCATTA TTTGCTAGCC AGTGTAAATT CAAGTGTCAT	16920
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TGTAGCTCTG CTCTTGCAG GTTTGGGAAT CCAATCTTCT GTAGCAGGAG TTCCGTCTAA	17280
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CATCGTATCG GTTCTATTAG CTATTGTCAAT CCTTTACAAT CTGACCAATA TCAACGTAGC	17940

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 TTATCCGCAG GTAGGCTGGG AAGTCTATGT AATCCCAGTG GCAGCAGTAA GCATCATTAA 18180
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 GAAATCTGTA GAGTAAGGTA GTTATTTTA GCTGATTGAA CTTCTATTTA CTAATATTCA 18300
 AAAATCCTCC GTTCAAAGA GCAGGGAACT CTTTGTGACA GAGGATTTT TCTATAGGGC 18360
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 TTCGTTGATC AAGAGGGCAT AATCGCGCCC GAAAGAATGG TCAAAGTAGT CTGAAAGCAT 18540
 AATGGCATTG TCAAGGCCTT CAGCACCGCA CCAACGTTT TGAGCAAAG GTAGGTCCAT 18600
 TGAAACAGTC AATACGACCG TGTGTCCAG TCCAGCCAAT TCTTCATTAA AACGACGTGT 18660
 TTGAGTTGAG CAGATGCCCTG TATCGATAGA AGGAACGACA CTCAAGACTT TTTTCTTGCC 18720
 ATCAAAATCA GCCAGAGATT TTTAGAAAG ATCTGTTGTA GTAAGAGAAA AATCAAGCGC 18780
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 GTTCCCTCTT CTTCAAGTATT GATTTCCCAG ATGTAAGCAC TTGGGACACC ATAGGCTTTA 19260
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 TTAAATTTAT CCTTACTTTT TTTATCAAGT TTGCTTAAC TATCACTATA TTCTTTGAGA 19380
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 TTAGCAAAAA TAATACCGTT TTCAAGGTTA AGCCAAGCGT GTGGTCTTC TTTTCCCTTT 19500
 TCATTTGAC CTTCAAGGTA GATAACATCA ACAGCCGTGCG TGACTGCGAA GTAGTCTTG 19560
 TTTTCAGTTT TCTTGGCATT TTCTACCAAT TTTGTAAACC AAGCATTGCC ACCTGTTCA 19620
 AGGTTGATAC CGTTATAGAA AATCAAATTA GCCTCAGAAG TTTTCTTAAC GTCTTCAGGA 19680

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ACTGGTTCGT ATTCTGTTGG GTCTTGCCC ACGAATTCAGCA TACTATGAAG GTCAATTTC	19740
TCACCAAGCAA TATTTTTAGT AATATCAGCG ATGATTGAGT TTGATGCAAC AACCTTTAGT	19800
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AGAAAAGAGAA CGAGTAATGT ACCTAATTTT TTCATTAGAT CCTCCAATTT ATTAGGGCTT	19920
TGCCCTTAT TTTAACAAAT GTTTATTTT CAGTTCAAA TATCGTTGTT TGGGAGCGAT	19980
AAAGAAGCTA ATGAGAAAGA AACTAGCAGC TGTAAGCAGC ATACTAGAAC CTGCCGCAAC	20040
ATTAAGAACTA TAGCCAATAA AGAGTCCAA AACTGAAGCA GTAGCTCCGA AGGTTGAGGA	20100
AAGGAAAATC ATACTTTCA GACTATTAGC ATACAGATAA GCAGTTGCAG CTGGGGTAAT	20160
CAGCATGGCT ACAATCAGGA TAGTTCCGAC ACTTTGCATG GCTGTCACAG ACACGAGAGT	20220
CAGGAGTACC ATGAGAAGGT ATGTATGAA ATTGACAGGC ATTCCCATGG CTTTAGCCAA	20280
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AGCTGCCCAACACCCATAG TAATAAACAT ATCCGTATCT TGGACGGCCA GGATATTACC	20400
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TTTGATGTTAG GTAATGATGA TGGCAGCTAG CAATCCAAAG ACAATGGCTC CGATAAAGAA	20580
GTCAAGGCCCAAGATGAAGG ATAGGGCTAC ACCTGGTAAG ACAGCATGTG AAATGGCCTC	20640
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GACGACAATA GCTGTTATCA AGGCATTTTG TAGGAAATGG AATTTTGCA ATCCATCGAT	20760
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GTCTTCCAG CTTTTTCAA ATCTCTCAGC GTATTCTGATG TGATTTCTC ACTGACAGAG	21000
TCAATCCAG CAAAGGGTTC ATCCAAGAGG ATATAGTCGG CTTCCCTGCAC CAAACATCTG	21060
GCAATCAAGA CCCGCTGGAA TTGACCTCCA GACAGTTGAC TAATTTGACG TTCAGCGTAG	21120
TCAGCTAGGC CGACGATTTTC AAGGGCCTCT TGCACTTCTC TCCAATGTTT AGCCTTTAAA	21180
CTTCGAAAGA GAGGAATAGA GGGAAATAGT CCTAACGAGA CGCATTCTT GACCTTGATG	21240
GGAAAGTTGT AGTCGATATT GATTTTTGTT TCCACATAGG CAATTGGTG TAAGGATTTT	21300
TTAACCTCCT TGTCATCGAG AAATGCCCTGA CCTTGATGTG GGATAATTCC CAACATACCT	21360
TTTAATAGTG TTGATTTCCC AGCGCCGTTT GGACCAATGA TGCCGGTAAT TGTTGGTCCA	21420
TGGAGCACTA GTGAAATATC CTTAAGTGCC AACGTTCTT TGTAGGAGAC ACTGAGGTTT	21480

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TCGATACGTA TCATAAACTT GTATTCCTCC TGTCTCTTAA TATACATTAA AAAAAAAATT	21540
AAGTCAAGTT AATTTTGAA AAAATTAAA TAATAACTGA AAAATAGATT CTAAAGATAA	21600
CTTTCAGGAT AAATTCTAA ATTATAAAAC GCATAGTATC AAGTGTAAAA AACTTGGAAAT	21660
TATGCCGTTT ATCATGGAAA GATTTTTAT AATAGCTAAA AAATAA	21706

(2) INFORMATION FOR SEQ ID NO: 37:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 6171 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 37:

GATCCCCAGG AAAAACCGAG GTTTTCCCAA TCAATCGTTA CTGTCATATT CCACTCCTTA	60
TTCTAAAAAC CTATTTCTTA TATTCTACAC TATTTTCTA AAATAGCAAG TATATTTGT	120
AATTTCAAGA AAATTCTCC AATAAAAACC AACTCTTAGA ACTGATTCTT CATTTCACTT	180
ATTTATCTTC AGTAACTACT CCCTGAAGAT AACCGTCAAA AACTTCTTCA TCTGAAATCG	240
TGTCAGAAAT GAAGCTTCCA TTGCTAGTGC GTTCTGACAA GTTCAAGTCT TGCAATCGGC	300
TTTCATAGAT TGTTCTTTA TTGGATTGGA CAAGCAGAGT TTGGTCGTTACATCCACTT	360
CCGTACTCMA GAAATGCCA ACAAAATCCIT GCTCTGCAAC TGCTCCTGCC AAGAAGACAC	420
GATGCCGTTT GTTTTCAAC TCACGCAAGA CTTGTAATCC TCGTTGGCA CGGCTGGTTG	480
CTAGAATTTC CTCAATGGAA ACACGTTCA AGCTTCCACG CTGGGTCAAG AGGTAGAAGG	540
ACGAAGTATT ACAGATAAAAG CCAGATTGGA GGACATCATC TTCTTCAAA TTCATAGCCT	600
TGACACCTGC TGCCCTTAGCA CCGACAACCG GAAACCTCTTC GATATTGAAA CGCAGGGCAT	660
AACCATTTC ACTAACCAAG ACAACATCAT CTAGTTTAAT CGGAGCCACT GCTACAATCT	720
GATCTGTATC GTCTTGAGC TTAGCATACT TGACAGACTT AGATCTATAG GTCCGCCATG	780
GAGTGAATTTC TTTTCGCTCT ACCCGTTGA TTGACCAAG GCGAGTCACT GCAAAGTAGG	840
TTGTCGCATC GTCAAACGTGA TCCAGTACTT CCACATAAG GATTCTTCA TTCGTTCAA	900
AGTTTGAT GGTTGGCTC AGATGCTCTC CGATGTCCTT CCAACGAATA TCTGCCAACT	960
CATGGATTGG TCTGTAGATG ACATTCCAA GACTTGTGAA CATCAAGAGG TGCTGGTTG	1020
TCTTGGCAGA TTGAAACAAA ATCAAACGGT CATCATCACG CTTGCCAATT TCTTCCAAGG	1080
TGGAAGCCGC AAAGGAACGT GGACTGGTAC GCTTGATGTA ACCTGCCTTG GTCACGCTGA	1140

372	
CGTAGGTATC TTCCCTCAGCG ATAAGACTAG CTGTATCAAT CTCATTGCT TTCGCAGTGT	1200
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TCATGAGATT GTACATAGTC CTTCATCAC CGATAATAGC CGCCAGCATA GCAATCTTCT	1320
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GTTGCAAAGT TACGATAGCC TCAGCCTGTT CTTCCGTAAA ATCATAGCTA ACTTGAGGT	1440
TTTCTTGGC GTCCGCCCTTA TTCTCAGAAG CACGGATAAG AGCAATGACT TCATCCAAA	1500
TCGAAATCAC ACGAATCAA CCTTCGACGA TATGGAGACG TTTCTCAGCC TTTTCTTGT	1560
CAAAGCGTGA ACGGCCAAA ATCACTTCTC GACGGTGAGC GATATAGCTA GACAGGATTG	1620
GAACAATCCC AACCTGACGA GGTGTGAAAT TGTCAATCGC CACCATATTA AAGTTGTAGT	1680
TGATTGTAG GTCGGTGTAC TTAATAAGT AGTTGAGAAC AAGCTCAGTA TTAGCGTCTT	1740
TCTTAAGTTC GATAGCGATA CGAAGACCAT CACGGTCAGA CTCATCACGA ACCTCAGCAA	1800
TCCCAGCTAC CTTGTTATTA ACACGAACAT CATCGATTTT CTTGACTAGA TTGGCCTTAT	1860
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TTTCAGTCTT GGAACGAACA ACCACGCGCC CTTTCCCACT CTCATAAGCT TTCTTGATT	1980
CATCACGACC CTGAATAATA GCCCCTGTAG GGAAGTCTGG TCCAGGCAAG AATTCCATGA	2040
GTTTATCAAT CTTTGCAGTT GGGTGGTCAA TCATGTAAAC TGAGCATCT ATGACCTCAG	2100
CTAAATATG GGGAGGAATG TCTGTGGCAT AACCAAGCCGA AATCCCAGTC GAACCATTGA	2160
CCAAGAGTT TGGAAAGGCT GCTGGCAAGA CCGTTGGTTC TTTCTCCGTA TCGTCAAAGT	2220
TCCATGCAA AGGAACGTGTC TTTTCTCGA TATCCTGAAG AAGGTAGCCT GCAATTTCAG	2280
ACAAACGTGC CTCAGTATAA CGCATAGCCG CAGGAGGATC TCCGTCCATA GAACCGTTAT	2340
TACCGTGCAT TTCAACTAGA ATCTCACGAT TTTTCCAGTT CTGTGACATA CGAACCATGG	2400
CATCATAGAT AGAAGAATCC CCGTGTGGGT GGAATTCCC CATGATGTT CCGACTGACT	2460
TGGCCGACTT ACGGTAGCTC TTGTCAAAAG TATTGCTATC CTTATTTCATA GAATAAAGAA	2520
TACGGCGCTG AACCGGCTTC AACCCATCAC GAATATCTGG CAAAGCCCG TCTTGAATAA	2580
TGTACTTGGA GTAGCGACCA AAGCGCTCTC CCATGATGTC CTCCAGGGAC ATGTTTGAA	2640
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TTTTCCAAAA TTAGTCTTAG TTTGTGTCTT AGCCGCTCCC TTAAGCGCCT CTTTGAGATA	2940

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AGCACTCATA GCAGATTCTT CATTAATAAT CCTGCAATT TTTCAAACCA AGATTTCAA	3000
ACTGCCTTTT CACATAGTC TTCACATCCG ACTCTAATT CCAGTTACT AACATATTAT	3060
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TAAATCTTC ATTGTGACAC GGATGAGGGT ACGTGTTCT GGGTTCATGG TTGTTTCCC	3240
GAGCTGGTCC GCATTCATCT CACCAAGTCC TTTGTATCGT TGGAGGGTAG CGCCTTTACC	3300
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CTTGCCTTTA CCTTGGACA TCTTGTAAAG AGGTGGGAGG GCAATATAGA CATGACCTGC	3420
CTCGACTAGC GGACGCATGT AACGGTAGAA AAATGTCAAG AGCAAGGTCT GGATATGGC	3480
ACCGTCGGTA TCCGCATCGG TCATGATAAT GATCTTATCA TAGTTGGCAT CTTCAATAGA	3540
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AGCCTGGAAC TTGGGTACAC GACCTGTTT GGCAGAACCA CCGCAGAGT CCCCCCTAAC	3720
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CAAGCCCTTA TCTTCTTGT TTTTCTTCCC ATTTCGGCTC TCATCACGCC CTTACGTGC	3840
TGCTTCACGA GCATCACGGG CCTTGATAGC CTTGGGATG AGGTTAGAAG CTAATTCCCC	3900
ATTTTCCATA AGGAAAAAAGG TCAACTTATC AGCCACTATT CCATCCACAA CTGGCGAGC	3960
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CTATTCTACA GGTTTCCAA GGATTTGCA AAATTTTCT TTCTCCGATG TGACAATTTC	5100
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GTTTATTAA TCCTAGCCTA TCTGCTGGGT TCGATTCCAT CTGGTCTCTG GATTGGACAA	5220
GTATTCTTC AAATCAATCT ACGCGAGCAT GGTTCTGGTA ACACTGGAAC GACCAACACCC	5280
TTCCGCATTT TAGGTAAGAA AGCTGGTATG GCAACCTTTG TGATTGACTT TTTCAAAGGA	5340
ACCCTAGCAA CGCTGCTTCC GATTATTTT CATCTACAAG GCGTTCTCC TCTCATCTTT	5400
GGACTTTGG CTGTTATCGG CCATACCTTC CCTATCTTG CAGGATTAA AGGTGGTAAG	5460
GCTGTCGCAA CCAGTGTGG AGTGTATTC GGATTTGCGC CTATCTCTG TCTCTACCTT	5520
GCGATTATCT TCTTGGAGC TCTCTATCTT GGAGTATGAA TTTCAGTGTCA TAGTGTACA	5580
GCATCGATTG CGGCTGTTAT CGGGGTTCTG CTCTTCCAC TTTTGGTTT TATCCTGAGT	5640
AACTATGACT CTCTCTTCAT CGCTATTATC TTAGCACTTG CTAGTTGAT TATCATTGCT	5700
CATAAGGACA ATATAGCTG TATCAAAAT AAAACTGAAA ATTTGGTCCC TTGGGGATTG	5760
AACCTAACCC ATCAAGATCC TAAAAAAATAA ATGCCAGTT CTGTAATGCC CCCAACAGT	5820
TAGACAAATA ATTTATCAA AGGATTTAGT TCTGTACTGC ACAGGACTAA GTCCCTTTAG	5880
TTTTACCTTA ATTGTTGT TGTGTAGTA ATCAATATAG TCTATAATGG CTTGTTCCAA	5940
TTGATTAAGT GATTAAATG TTTTCTCATA GCCATAAAAC ATTCGGATT TTAAATGCC	6000
AAAGAAAGAT TCCATCCTAC CGTTGTCTTG GCTGTTGCC TTACGTGACA TGGATGCTTG	6060
AATTCCCTTA CTCTCTAGGA ACCGATGATA AGAATCGTGT TGGTATTGCC AGCCTGGTC	6120
ACTATGGAGA ATCGTATTCT CGTAGTGCTT CTCTGTGAAT GCCTGTTCCA A	6171

(2) INFORMATION FOR SEQ ID NO: 38:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 18475 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 38:

TATTACAAAT	AAAAAACGG	AGGAGTGCTT	TATGAAAGCC	TATACTTATG	TTAAACCAGG	60
ACTTGCTCT	TTTGTGATG	TAGACAAACC	AGTTATTCGC	AAGCCAACAG	ACGCTATTGT	120
GCGTATTGTA	AAAACCACTA	TTTGTGGAAC	AGACCTCCAT	ATTATCAAAG	GGGATGTTCC	180
TACTTGCCAA	AGTGGTACCA	TTCTTGCCCA	CGAAGGGATT	GGGATTGTTG	AAGAAGTTGG	240
GGAAGGAGTT	TCCAACCTCA	AAAAAGGTGA	CAAGGTCTTG	ATTTCTTGCG	TCTGTGCCTG	300
TGGTAAATGC	TACTACTGTA	AAAAGGAAT	TTATGCTCAC	TGTGAAGACG	AAGGGGGCTG	360
GATTTTCGGT	CACTTGATTG	ATGGTATGCA	GGCTGAATAT	CTACGTGTCC	CTCATGCAGA	420
TAATACTCTT	TACCATACTC	CAGAAGACCT	GTCAGATGAA	GCTTTGGTTA	TGCTGTCAAGA	480
CATTCTGCCT	ACTGGATATG	AAATTGGTGT	CTTAAAAGGG	AAAGTAGAAC	CTGGTTGCAG	540
CGTAGGCCATT	ATGGGTTCAAG	GTCCAGTTGG	ATTGGCTGCT	CTTTAACAG	CCCAATTCTA	600
TTCACCAGCT	AAATTGATTA	TGGTAGACCT	AGACGATAAC	CGCTTGGAAA	CTGCCCTATC	660
ATTCGGTGCG	ACTCATAAGG	TTAATTCTTC	AGACCCCTGAA	AAAGCCATTA	AAGAAATTAA	720
TGATTGACA	GATGGTCGTG	GTGTGGATGT	CGCTATCGAA	GCTGTTGGTA	TTCCCTGCAAC	780
ATTTGATTTG	TGTCAAAAGA	TTATCGGTGT	AGACGGAACG	GTTGCCAACT	GTGGTGTGCA	840
TGGTAAACCA	GTTGAAATTG	ATTTAGATAA	ACTTTGGATT	CGCAACATCA	ATGTAACAAAC	900
TGGTTTGGTA	TCTACAAAATA	CGACTCCACA	ATTGTTGAAA	GCACTTGAAA	GTCATAAGAT	960
TGAACCGGAA	AAATTGGTAA	CTCACTATTT	CAAACTCAGT	GAAATTGAAA	AAGCCTACGA	1020
AGTCTTCAGT	AAGGCAGCAG	ACCAACATGC	CATTAAGGTC	ATTATCGAA	ACGATATCTC	1080
AGAAGCCTAA	GTAGTAAAAA	TATTTTGTA	CATAAGTAAA	TAGAAATTCA	GTCATCCATC	1140
AGATGGCTGG	ATTTTTTATC	AAAAAATTAA	GAAATGAGCA	TATTTCTTTC	CTTGCTCTGGC	1200
GGAATTGGTT	ATAATATACG	GTACAAAGGA	ATGAATGAAT	ATGTATCGTG	TTATAGAAAT	1260
GTACGGAGAT	TTTGAACCGT	GGTGGTTCTT	AGAAGGGTGG	GAAGAAGATA	TTGTAGCAAG	1320
TAGAAAATTAA	GACCACTATT	ATGATGCTCT	CAAATACAC	AAAACCTGCT	GGTTTAGATT	1380
GGAACAAGAA	TCGCCTCTTT	ATAAAAGTAG	AAGCGACTTG	ATGACCATT	TTTGGGACCC	1440
GGAAGACCAA	CGCTGGTGTG	ATGAATGTGA	TGAGTATTAA	CAACAATACC	ATTCTTTGGC	1500
TCTTTGCA	GATGAGCAGG	TTATCCCAGA	CGAAAAACTA	CGCTCAGGCT	ATGAAAAACA	1560
AACCAGTCAG	GAAAGGAATC	TTCTTGCCG	TATGAAATTAA	AAATAGAGAA	AAGTAACATT	1620
TTTGGAGTTG	CTTTTTTAT	TTTCTAACT	CTTTCGAAT	AGTATAGGTG	AGGAGGTAAG	1680

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TATGGTTCAA GAAATTGCAC AAGAAATCAT TCGTTCAGCT CGGAAAAAAG GGACGCAGGA	1740
TATCTATTTT GTCCCTAAGT TAGACGCCA TGAGCTTCAT ATGAGGGTAG GAGACGAGCG	1800
CTGTAAAATT GGTAGCTATG ATTTGAAAAA GTTTCAGCC GTTATCAGTC ACTTTAACGTT	1860
TGTGGCGGGT ATGAATGTGG GAGAAAAAAG ACGTAGTCAA CTGGGTTCCGT GTGATTATGC	1920
CTATGACCAT AAGATAGCGT CTCTACGTTT ATCTACTGTA GGCGATTATC GGGGCATGA	1980
GAGTTGGTT ATCCGTTGT TGACGATGA GGAGCAGGAC CTGCATTTTG GTTTCAGGA	2040
TATTGAAGAA TTAGGCAAGC AGTACAGGCA ACGGGGACTC TATCTTTTG CTGGTCCGGT	2100
TGGGAGTGGT AACACGACCT TGATGCATGA ATTGTCCAAG TCACCTTTTA AAGGACAGCA	2160
AGTTATGTCC ATCGAAGATC CTGTCGAAAT CAAGCAGGAC GACATGCTTC AGTTGCAGTT	2220
GAACGAAGCA ATCGGCCAA CCTATGAAAAA TCTAATCAA CTTTCCTTGC GTCATCGACC	2280
AGATCTCTTG ATTATCGGAG AAATTCGTGA CAGCCAGAGC GCGCGTGCAG TGGTCAGAGC	2340
TAGTTGACA GGTGCGACAG TCTTTCAAC CATTCACGCC AAGAGTATCC GAGGTGTTA	2400
TGAGCGTCTG CTGGAGTTGG GTGTGAGTGA AGAAGAATTG GCAGTTGTTG TGCAAGGAGT	2460
CTGCTACCAAG AGATTAATCG GGGGAGGAGG AATCGTTGAC TTTGCAAGCA GAGATTATCA	2520
AGAACACCAA GCAGCCAAGT GGAATGAGCA AATTGACCAAG CTTCTTAAAG ATGGACATAT	2580
CACAAGTCTT CAGGCTGAGA CGGAAAAAAT TAGCTACAGC TAAGAAAAA AATATCATCA	2640
CCCTATTTAA CAATCTCTTT TCTAGCGGTT TTCATCTGGT GGAGACTATC TCCCTTTAG	2700
ATAGGAGTGC TTTGTTGGAC AAGCAGTGTG TGACCCAGAT GCGGTGCGGC TTGTCAGG	2760
GGAAATCATT CTCAGAAATG ATGGAAAGTT TGGGATGTTA AAGTGCTATT GTCACTCAGT	2820
TATCCCTAGC TGAAGTTCAT GGCAATCTCC ACCTGAGTTT GGGAAAGATA GAAGAATATC	2880
TGGACAATCT GGCTAAGGTC AAGAAAAAAT TGATTGAAGT AGCGACCTAT CCCTTGATT	2940
TGCTGGTTT TCTCTCTTA ATTATGCTGG GGCTACGGAA TTACCTGCTC CCACAACCTGG	3000
ATAGTAGCAA TATTGCCACC CAAATTATCG GTAATCTGCC CCAAATTTTT CTAGGCATGG	3060
TAGGGCTTGT TTCCGTGCTT GCCCTTTAG CACTCACTTT TTATAAAAAGA AGTTCTAAGA	3120
TGAGTGTCTT TTCTATCTTA GCACGCCCTTC CCTTTATTGG AATCTTGAGT CAGACCTACT	3180
TGACAGCCTA TTATGCACGT GAATGGGGGA ATATGATTTAC ACAGGGAATG GAGTTGACGC	3240
AGATTTTCA AATGATGCAG GAACAAGGTT CCCAGCTCTT TAAAGAAGTC GGTCAAGATC	3300
TGGCTCAAAC CCTGAAAAAT GGCGTGAAT TTCTCAGAC GATAGGAACC TATCCTTCT	3360
TTAGGAAGGA ATTGAGTCTC ATCATAGAGT ATGGGAAAGT TAAAGTCCAAG CTGGGTAGTG	3420
AGTTGAAAT CTATGCTGAA AAAACTTGGG AAGCCTTTT TACCCGAGTC AACCGCACCA	3480

TGAATTTGGT GCAGCCACTG GTTTTTATCT TTGTGGCACT GATTATCGTT TTACTTTATG	3540
CGGCAATGCT CATGCCCATG TATCAAAATA TGGAGGTAAA TTTTTAAAAT GAAAAAAATG	3600
ATGACATTCT TGAAAAAAGC TAAGGTTAAA GCTTTACAT TGGTGGAGAT GTTGGTGGTC	3660
TTGCTGATTA TCACGCTGCT TTTCTTGCTC TTTGTACCTA ATCTGACCAA GCAAAAGAA	3720
GCAGTCAATG ACAAAAGAAA AGCAGCTGTT GTTAAGGTGG TGGAAAGCCA GGCAGAACTT	3780
TATAGCTTAG AAAAGAACATGA AGATGCTAGC CTAAGAAAGT TACAAGCAGA TGGACGCATC	3840
ACGGAAGAAC AGGCTAAAGC TTATAAAGAA TACAATGATA AAAATGGAGG AGCAAATCGT	3900
AAAGTCAATG ATTAAGGCCT TTACCATGCT GGAAAGTCTC TTGGTTTGAG GACTTGTGAG	3960
TATCCTTGCC TTGGGCTTGT CCCGGCTCTGT CCAGTCCACT TTTTCAGCGG TAGAGGAACA	4020
GATTTCTTT ATGGAGTTTG AAGAACTCTA TCGGGAAACC CAAAAACGCA GTGTAGCCAG	4080
TCAGCAAAAG ACTAGTCTGA ACTTAGATGG GCAGACGCTT AGCAATGGCA GTCAAAAGTT	4140
GCCAGTC CCT AAAGGAATTC AGGCCCCATC AGGCCAAAGT ATTACATTG ACCGAGCTGG	4200
GGGCAATTG TCCCTGGCTA AGGTTGAATT TCAGACCACT AAAGGAGCGA TTGGCTATCA	4260
ATTATATCTA GGAAATGGAA AAATTAAACG CATTAAAGGAA ACAAAAATT AGGGCAGTGA	4320
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GACAAATMCA AAAAATAGG CAAGAGGAAG CAAAATCTT GCAAAAGGAA GAACTCTTGA	4440
CCCTAGCTAA GATGCCCTG CAGACGGGGC AAAATCAGGT AAGCATCAAC GGAGTTGAGA	4500
TTCAGGTATT TTCTAGTGA AAAGGATTGG AGGTCTACCA TGGTTCAGAA CAGTTGTTGG	4560
CAATCAAAGA GCCATAAGGT CAAGGCTTT ACCTTGTAG AATCCCTGCT TGCCCTCAT	4620
GTCATCAGTG GGGGATTACT CCTTTTCAA GCTATGAGTC AGCTCCTCAT TTCAGAAGTT	4680
CGCTACCAGC AACAAAGCGA GCAAAAGGAG TGGCTCTTGT TTGTGGACCA ACTTGAGGTA	4740
GAATTAGACC GTTCGCAGTT CGAAAAAGTA GAAGGCAATC GCCTATACAT GAAGCAAGAT	4800
GGCAAGGACA TCGCCATCGG TAAGTCAGG TCAGATGATT TCCGTAAAAC GAATGCTCGT	4860
GGTCGAGGTT ATCAGCTAT GCTTATGGA CTCAAATCTG TACGGATTAC AGAGGACAAT	4920
CAACTGGTTC GCTTCATTT CCAGTTCCAA AAAGGCTTAG AAAGGGAGTT CATCTATCCT	4980
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CTTTAGTCTT TTGTTGCAAT TTTATTTGAA CCGACAAGTC GCCCACTATC AAGACTATGC	5100
TTTGAATAAA GAAAAATTGG TTGCTTTGCA TATGGCTAAA CGAACCAAAG ATAAGGTTGA	5160
GCAAGAAAGT GGGGAACAGT TTTTAATCT AGGTCAAGTA AGCTATCAA ACAAGAAAAC	5220

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TGGCTTAGTG ACGAGGGTTC GTACGGATAA GAGCCAATAT GAGTTTCTGT TTCCCTTCAGT	5280
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AGTGGAGAAG AAAAATCAG AAGAGAAGCC TGAAAAGAAA GAGAATTCAT AGTCAATTCA	5400
ACTATAATGC GTTGAATCCA GAATAGTCCA CTGTAGTTTC TAGAAAATTG CTGGAAATGG	5460
ATGTTAAGCT CCAATTCAATT TGTATTATAC TTATTCAGT TTACTATACT TTGTGCTAAA	5520
TTAAAGATAT GAAACATGAT TTTAACACAA AAGCAGAAC TTTCGATTCC CCTAAAAATA	5580
TCTTCCTCGC AAACTTGGTA TGTCAAGCAG CCGAGAAACA GATTGATCTT CTATCAGACA	5640
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GACCTGTTTC AAGGAAATCA CTCAGAATTTC TTTTAATAG TAGCCAAAA ATCACTCGCC	6120
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GCACTACGCA AAGAAGAATG GCTCAAGACC TACCAAGTTTC TCTTGATGAA GGCTGGGCAA	6420
ACAGAACCCCT TGCAGGCCAA TCACCAGTTT ACACCGGATG CTATTGCTTT GCTTTGGTG	6480
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GTTGCTTCTA GCCAAGAACAA TACTTACGCC CATCACTTGC TCATGAAACA AGGGCTTAAG	6840
TACCTCAACT CAGACGGATA CGCTATTTTCTAGCTCCGA GTGATTTGTT GACCAGTCCT	6900
CAAAGTGATT TGTAAAAAGA ATGGCTGAAA GAAGAGGCCA GTCTGGTTGC TATGATTAGT	6960
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AAAAATGAAA TAGCAGTAGA GCCTTTGTT TATCCACTTG CTAGCTTGCA AGATCCAAGT	7080
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AGATTTGTT ATAATAGTTG AAAACGCTTA AAAAGGGTA TCATGTTATG ACAAAAACAA	7200
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AAATTTTATT GGATGACTTG ATTGTTTCG ATATTATCAA GGCTTATGAC GAGATTACAG	7440
GTGTTGGACA TCGTGTGTT GCTGGTGGAG AATATTCAA AGAACATCAACA GTTGTGAGG	7500
GAGATGTTT AGAAAAAGTT GAAGAGTTGA GTTTGTTGCC TCCTCTACAC AACCCGGCCA	7560
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GTCATATTGG TAACGGAGGC TCAATTACAG CTGTGAAAGC CGGCAAATCT GTAGACACTT	7860
CTATGGGTT CACTCCTCTT GGTGGTATTA TGATGGAAC GCGTACAGGG GATATTGATC	7920
CAGCTATCAT TCCTTATTTA ATCCAATATA CAGAGGATTT TAACACACCA GAAGATATCA	7980
GTCGTGTCT TAACCGTGAA TCAGGTCTTT TGGGAGTTTC TGCTAATTCT AGCGATATGC	8040
GCGATATAGA AGCAGCTGTA GCAGAAGGGATCACGAGGCTAGCTTGCT TATGAAATGT	8100
ATGTTGACCG TATCCAAAAA CATATCGTC ACTACCTTGC AGTGCTAAAT GGAGCAGATG	8160
CCATTGTTT CACAGCAGGT GTCGGTGAAA ATGCAGAGAG TTTCCGTCGT GATGTAATCT	8220
CAGGGATTTC GTGGTTGGT TGTGATGTTG ATGATGAAA GAATGTCTTT GGCGTTACAG	8280
GAGACATCTC AACAGAGGCC GCTAAAATCC GTGTCTTGGT TATTCCAACA GATGAAGAAC	8340
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TACAAGGAGT TCGGAAAGTT ATTTTCCAG CTTCTTTTC TGATGAAATT GTCCAAAACC	8460
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ATTAAGCGAC GGATTCGGCA TATTATCCAG AATGCAAAG GGAGTCTGGT AGAAGATGTC	8760

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GAAACTAAAG	TTGACTAGTT	TGCTAGGACT	GTCTCTGTTA	ATCATGACAG	CCTGTGCCAC	8940
TAATGGGGTA	ACTAGCGATA	TTACAGCCGA	ATCGGCTGAT	TTTTGGAGTA	AATTGGTTA	9000
CTTCTTTGCG	GAAATCATTG	GCTTTTATC	GTTTGATATT	AGTATCGGAG	TGGGGATTAT	9060
TCTCTTACG	GTCTTGATTC	GTACAGTCCT	CTTGCCAGTC	TTTCAGGTGC	AAATGGTGGC	9120
TTCTAGGAAA	ATGCAGGAAG	CTCAGCCACG	CATTAAGGCG	CTTCGAGAAC	AATATCCAGG	9180
TCGAGATATG	GAAAGCAGAA	CCAAACTAGA	GCAGGAAATG	CGTAAAGTAT	TTAAAGAAAT	9240
GGGTGTCAGA	CAGTCAGACT	CTCTTGCCC	GATTTGATT	CAGATGCCGG	TTATTTGGC	9300
CCTGTTCCAA	GCCCTATCAA	GAGTTGACTT	TTTAAAGACA	GGTCATTTCT	TATGGATTAA	9360
CCTTGGTAGT	GTGGATACAA	CCCTTGTTCT	TCCGATTTA	GCAGCAGTAT	TCACCTTTT	9420
AAGTACTTGG	TTGTCNAACA	AAGCTTGTCT	TGAGCGAAAT	GGCCTACGA	CTCGATGAT	9480
GTATGGGATT	CCAGTCTTGA	TTTTTATCTT	TGCAGTTAT	GCGCCAGGTG	GAGTCGCCCT	9540
ATACTGGACA	GTGCTAATG	CTTATCAAGT	CTTGCAAACC	TATTCTTGA	ATAATCCATT	9600
CAAGATTATC	GCAGAGCGCG	AGGCCGTAGT	ACAGGCACAA	AAAGATTTGG	AAAATAGAAA	9660
AAGAAAAGCC	AAGAAAAAGG	CTCAGAAAAC	GAAATAAATA	AGGAGGAATC	TGGTAGTGGT	9720
AGTATTTACA	GGTCAACTG	TTGAAGAAGC	AATCCAGAAA	GGATTGAAAC	AATTAGATAT	9780
TCCAAGAATG	AAGGCTCAT	TCAAAGTCAT	TTCTAGGGG	AAAAAAGCT	TTCTTGGTCT	9840
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GAAGACGGTT	AGTGAAGAAA	CCGTTGACCT	TGGTCATGTG	GTTGATGCTA	TTAAAAAAAT	10020
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ACATGCCAGC	ACTATCTTAG	AAGAAACTGG	TCACATTGAG	ATTTTAAATG	AACTTCAAAT	10140
CGAGGAAGCG	ATGAGGGAAG	AAGCAGGCCGC	TGATGACCTT	GAAGACTGAGC	AAGACCAAGC	10200
TGAAAGTCAA	GAACTAGAAG	ACTTGGCTT	GAAAGTTGAA	ACGAACCTTG	ATATTGAACA	10260
AGTAGCTACG	GAAGTAATGG	CTTATGTTCA	AACGATTATT	GATGACATGG	ATGTTGAGGC	10320
TACACTTICA	AATGATTATA	ACCGTCGTAG	CATCAATCTA	CAAATTGACA	CCAACGAACC	10380
AGGTCGTATT	ATCGGCTACC	ATGGTAAAGT	CTTGAAGGCC	TTGCAACTGT	TGGCTAAAAA	10440
TTATCTTAC	AACCGCTATT	CCAGAACCTT	CTACGTTACA	ATCAATGTCA	ATGATTATGT	10500
CGAACACCGT	GCAGAAGTCT	TGCAAGACCTA	TGCGCAAAAA	TTGGCGACTC	GTGTTTGGGA	10560

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GAGGTTAACAC TGATGTTGAA TAAGATAAGA GACTATTTAG ACTTTGCTGG TTTGCAGTAC	10800
CGTAACTCCTG ATAAAGCGGG AGCAGAGCGA GAGAAGATGC TGGCATTCCG CCACAAAGGA	10860
CAAGAGGCC GAAAGGTTTT TACAGAACTG GCCAAAGCCT TTCAAGCAAG CCATCCAGAA	10920
TGGCAACTCC AACAGACTAG CCAGTGGATG AATCAGGCC AGCGTTGAG ACCACATT	10980
TGGGTTTATC TACAGAGAGA CGGACAAGTG ACAGAACCTA TGATGCCCTT ACGTTTGAT	11040
GGGACATCTA CTGACTTGG AATTTCTTG GAAGTCAGTT TCATCGAACG TAAGAAGGAT	11100
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CCTATGACAG AAAATTGTC TGAAGAAGAA ATCGTAGAAG GCTTTATTGAA GTCTTATTCT	11340
AAAATTCTTC CCTATTATCT AGCTACGAGA AAATAAGATA ATTTGTAAAA CATCATAAAT	11400
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GCATTATATA ATAAACTTAC AAAAACAAATT CAAAAGGAGA ACAATTATGG AAGTCGTTTC	11520
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GGCAACTCAA CGCTTGACTG GTGGTGGCGG ATTTGCGATT GGTCACCAAC AGCAATTGCG	12120
AATCTGGTT GTAGATAAAAG TAGCAGGACG CTTGGTAAG AAAGAAGAAA GTTTAGACAA	12180
TCTTAAATTA CCTAAGTTCC TCTCAATCTT CCACGATACA GTTGTGTCAT CTGCTACCTT	12240
GATGCTCGTA TTCTTCGGAG CCATTCTTTT AATCTTGGGT CCAGACATTA TGTCTAATAA	12300

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AGAAGTCATC ACTTCAGGAA CTCTATTCAA TCCTGCTAAA CAAGATTTCT TTATGTACAT	12360
TATCCAAACA GCCTTACCT TCTCAGTTA CTTGTTCGTT TTGATGCAAG GTGTCCGAAT	12420
GTTCGTATCT GAGTGACAA ACCGCTTCCA AGGTATTCA AACAAATTGT TGCCAGGTT	12480
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GGTCTACGCT GATAAACGCG GCGGATGGAA ACGGGCTGTT ATCCTTCCT TTATATCAGG	12720
TGTCCTTCAA GTTGCTCTAG GAGCTTTTG TGTGGCCCTT CTCGATTTGG CATCTTATGG	12780
TGGCTACCAT GGAAATATCG ACTTTGAATT CCCATGGCTT GGATTTGGAT ATATCTTCAA	12840
ATACCTTGGT ATTGTTGGTT ATGTAATTGT GTGCTCTTC TTGCTTGTAA TTCCCTCAACT	12900
TCAATTGCCC AAAGCAAAAG ATAAAAGAGAA ATATTACAC GGTGAAGTTC AAGAAGAAC	12960
TTAGTATCTA GAAAAGGAGA AATAAAATGG TAAAGTATT AGCAGCGTGC GGAAATGGAA	13020
TGGGTTCATC AATGGTTATC AAGATGAAGG TTGAAAATGC TCTCCGTAAG CTTAATCAA	13080
CAGATTTAC AGTCAATTCA TGCACTGTG GTGAAGCTAA AGGTTTAGCA GTAGGATATG	13140
ACATCGTAAT CGCTCTCTT CATTGATTC AAGAATTGGA AGGGCGAACT AATGGGAAGT	13200
TAATTGGGCT TGATAACTTG ATGGATGATA AAGAAATCAC CGAAAAACTC AGTCAAGCAC	13260
TACAGTAAAA GGTTGGAGGG GGCTGGACAG AAACGTGAGAG TTATCGTTTC TGTCCTTCTC	13320
CCTCTTAAA TAAAGGGAGC AGATATGAAT TAAACAAAG CTTTAATTGA CAATGACTCG	13380
ATCCGACTAG GTTTAGAGGC TAACAATTGG AAAGAAGCAG TCAAGGTAGC AGTAGATCCC	13440
TTAATTGAAA GTGGGGCAAT TTTGCCAGAG TATTACGATG CTATCATTGA ATCGACTGAA	13500
GAGTATGGGC CTTACTATAT CTTGATGCCA GGTATGGCTA TGCCCCACGC TAGACCTGAA	13560
GCAGGTGTGC AAAGTGTGTC CTTTCATTG ATTACCTTAC AAAATCCTGT TGTATTTCA	13620
GATGGAAAG AGGTATCTGT TTTGTTGGCA CTAGCAGCAA CAAGTTCAA AATTACACACA	13680
AGTGTAGCCA TTCCACAAAT TATTGCCCTA TTGAAATTAG AAGATTCTAT TGCACGTTA	13740
CAGGCTTGCC AGACTAAAGA AGATGTCTG GCTATGATTG AAGAATCTAA GGATAGCCCT	13800
TATCTCGAAG GATTGGATTG GGAAAGTTAG AAAGAGGAAT AAAGAAATGA CAAAAAGAAT	13860
ACCTAAATTAA CAAGTTGCAT TAGACCATTG AGACTTGCAA GGAGCGATTAA AAGCAGCTGT	13920
TTCTGTTGGT CAGGAAGTAG ATATTATCGA AGCTGGAACG GTTGCTTGTG TTCAAGTTGG	13980
AAGTGAAC TG GCTGAAGTCT TGGTAGCCT TTTCCAGAT AAGATTATTG TGGCAGACAC	14040
AAAATGTGCT GATGCTGGTG GAACAGTTGC TAAAAATAAT CGCGTTCGTG GAGCAGACTG	14100

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GATGACTTGT ATCTGTTGTG CAACCATCCC TACTATGGAA GCAGCTCTAA AGGCTATCAA	14160
GAATGAAACGA GGAGAACGAG GCGAAATCCA GATCGAGCTT TATGGCGATT GGACTTTGA	14220
ACAAGCTCAG CTTGGCTAG ATGCAGGTAT CTCACAAGCT ATTATTCACC AATCTCGTGA	14280
TGCTCTCTT GCTGGTGAAA CTTGGGTGA AAAAGACCTT AATAAGGTTA AAAAACTCAT	14340
TGACATGGGC TTCCGTGTAT CTGTAACAGG TGGTCTAGAT GTAGATACTC TCAAACCTTT	14400
TGAAGGTATT GATGTCTTTA CCTTTATCGC AGGTCTGGGA ATTACAGAGG CTGTGGATCC	14460
AGCAGGAGCA GCGCGTGCCT TCAAGGATGA AATCAAACGA ATTGGGGGT AAATCATGGT	14520
ACGTCCAATT GGAATTTATG AAAAGGCAAC CCCAACACAC TGTACTTGGC TAGAACGTTT	14580
AAATTTGCC AAGGAGTTAG GCTTTGATTT TGTCGAGATG TCTATTGACG AACGTGACGA	14640
GCCTTAGCA AGACTTGACT GGAGTAAGGA AGAACGCTTG GAAGTTGTCA AAGCAATCTA	14700
TGAAACTGGT GTTCGTATTCT CTTCTATCG TTTTCAGGC CATCGTCGCT ACCCATTGGG	14760
TTCAAAAGAT CCAGTTCTAG AGGAAAAATC TCTAGAACTC ATGAAAAAAAT GTATCGAATT	14820
AGCTCAAGAC TTGGGAGTTC GTACGATTCA ATTAGCTGGT TACGATGTTT ACTATGAGGA	14880
AAAGTCACCC CAGACACGCC AACGTTTTAT CAAAAATTG AGAAAAGCCT GTGACTGGGC	14940
TGAAGAAGCT CAGGTGGTAC TTGCTATTGA ATTATGGAT GATCCTTTCA TCAGTAGCAT	15000
CGAAAAATAT TTGCTATAG AAAAAGAGAT TGACTCTCCC TTCCCTTTG TATATCCAGA	15060
TATTGGTAAT GTGTCTGCAT CGCATAATGA TATCTATAGT GAGTTTTATC TTGGTCATCA	15120
TGCCATCGCA GCTCTCCATC TCAAGGATAC TTATGCAGTG ACAGAAAGTT CAAAGGGCCA	15180
GTTCCGAGAT GTACCTTCG GGCAAGGTTG TGTCAAATGG GAAGAAGCTT TCGATATTTC	15240
AAAGGAAACC AATTATAATG GACCTTTCTT AATCGAAATG TGGTCTGAAA ATTGTGAAAC	15300
AGTAGAAGAA ACACGCGCAG CCATTCAAGA GGCGCAAGCT TTTCTCTATC CACTCATTAA	15360
GAAAGCAGGT TTGATGTAAG ATGAATCAAG TAATCAATGC TATGCGTAAA CGAGTCTGTG	15420
ATGCCAATCA ATCATTGCCA AAACATGGAC TTGCTAAATT TACCTGGGG AATGTATCTG	15480
AAGTTAATCG CGAAACTCGGT GTCATTGTTA TCAAACCATC AGGCCTGGAT TATGACGAAT	15540
TGACACCTGA AAACATGGTA GTGACTGATC TAGATGGTAA GATCCTAGAA GGGGATTTAA	15600
GACCACCTTC CGACCTCCCCA ACTCATGTGC AATTATATAA GACTTGGTCA GAAATTGGTA	15660
GTGTGGTTCA CACCCATTG ACAGAAGCTG TTGGTTGGGC TCAGGCAGGT CGTGATATTG	15720
CTTTCTACGG AACAAACCAT GCAGATTATT TCTACGGTTC AATCCCTTGC GCCCGTAGTT	15780
TGACCAAGGA CGAAGTAGAA GTGGCCTATG AAAAAGATAC TGGCCTGGTT ATCGTAGAAG	15840

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AGTTTGAACA TCGCGGACTT AACCCGGTTG AAGTACCAAGG AATTGTTGTA CGCAATCACG	15900
GTCCATTCACTCTGGGCAAA AATCCAGAGA ATGCTGTTA TCACCTGTC GTACTAGAGG	15960
AAGTATCAAA GATGAATCGC TTACAGAAC AAATCAATCC AAGAGTTGGA CCTGCTCCCC	16020
AGTACATACT AGAAAAACAC TACCAACGTA AACATGGACC AAATGCTTAT TATGGTCAA	16080
AGTAAGAACG ATGAAGGAGG AGAAAAAGAT AAATTTAGCT CCTCTTTTA CATTGATT	16140
TTATTGAGAG TAAAGTTGGA GTTGAAGTAA TTTTAAAAGA TTTTTTAGAA ATAGCGCTTG	16200
ATATATATAT GGTAAAATAA AAAGAATTGC TGTGATATCA ATAGATTGG GGGATTTTT	16260
AATATGGTAC TGGATAAGC AACGGTGTGAT TTGCTTCAAT ATTTGATGGA TCAAGAACG	16320
TCCAAAACGA TTATGGCGAT TTCGAAAGAT TTGAAAGAGT CAAGAAGGAA AATTTATTAT	16380
CACATTGACA AAATCAATGC TGCTCTGGGT GACGAGGCAC TTCACATCAT TAGTATTCCA	16440
CGAATTGGTA TTCACTTAAC GGAAGAGCG AGAGATGCTT GTTGTAAACT ATTATCGGAA	16500
GTAGATTTCGT ACGATTATAT CATGAGTGCG CATGAAACGTA TGATGATAAT GTTACTATGG	16560
ATAGGTATTT CAAAGAACG TATTACGATT GAAAATTGA TAGAGTTAAC AGAGGTATCT	16620
AGGAATACTG TTCTCAATGA TTTGAATAGT ATTGCTTATC AACTAACTTT GGAACAATAT	16680
CAGGTGATCT TGCAAGTGAG CAAGTCACAG GGATACAAACC TTCATGCCCA CCCTCTTAAT	16740
AAAATTCACT ATCTTCAATC GCTTCTATAT CATATTTTA TGGAAGAAAA TGCCACTTT	16800
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GAAATGAACC AATTTTTAA GGAACAGGTT CCTTTAGTTG AACAAGATTT AGGGAAAGAAA	16920
ATAAACCATC ATGAAATAAC TTTTATGTTG CAGGTTCTAC CTTATTTGCT GTTAAGCTGT	16980
CATAATGTTG AACAGTATCA AGAAAGACAT CAGGATATAG AGAAAAGATT TTCTTGATA	17040
AGAAAAAGAA TAGAGTATCA GGTGTCTAAG AAATTAGGAG AACGGTTGTT TCAAAAGTTT	17100
GAAATTCTT TGTCAGGACT TGAAGTTCTT CTTCTAGCTG TTCTCCTCCT CTCCATCGT	17160
AAAGATTTGG ATATTCAATC AGAAAAGTGAT GATTTTCGGC AATTTAAACT TGCTTTAG/A	17220
GAATTATCT GGTATTTGA ATCACAAATC CGAATGGAGA TTGAGAACAA GGATGATTTG	17280
TTACGAAATT TGATGATCCA CTGTAAGGCC TTGTTATTTA GAAAGACTTA CGGTATTTT	17340
TCTAAAAATC CTCTAACAAA ACAAAATCGA TCCAAGTATG GAGAATTATT TTTAGTCACT	17400
AGAAAAATCTG CGGAAATTAA AGAAGGAGCA TGGTTTATTC GGCTAACAGA CGATGATATT	17460
GCCTATTTGA CGATTCAAT TGGAGGATTG TAAATATA CACCATCATC TCAAAAAAAT	17520
ATGAAAAAAG TTTATCTCGT TTGAGTGAAG GGTGTTGCGG TTTCGAGACT TTTGCTGAA	17580
CAATGCAAAC TTTATTTCC AAATGAGCAA ATTGACACTG TATTACAAAC AGAACAAATT	17640

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AAGAGTGTGG AAGATATTGC ACAAGTTGAT GTAGTGATTA CTAACATGA TGATTTGGAT	17700
AGCAGATTC CGATTTAAG GGTAACTCCT ATCCCTGAAG CAGAAGATAT TTTGAAAATG	17760
CTAGACTATC TTAAACACAA TATATTCGT AATAAGAGCA AAAGTTCAAG TGAAAATCTT	17820
TCTAGCTTA TTTCGCTTA TATTGTAGAC AGCAAGTTGG CTAGTAAGTT CCAAGAAGAG	17880
GTTCAACAC TTATAAATCA AGAAATAGTA GTTCAAGCTT TTTTGAAGT TATTGAGG	17940
ACAGTCAAAT GATGAACACA AACCTGTGTk TTTCTGGTC TTTTGTAGTG TTTTGAAGGG	18000
TGGkATACTA ATCTCAAAGA TAACAATTAT ATCCAAGGA GGCAACATAT GCCAAACGTC	18060
AAAGAAATTA CAAGAGAGTC ATGGATTTA GCCACTTCC CAGACTGGGG AACATGGTG	18120
AACGAAGAAA TCGAAGAAGA AGTCGTACCT GAAGGCAACT TTGCCATGTG GTGGCTAGGC	18180
AACTGTGGTA CTTGGATTAA GACACCAGCT GGTGCTAACG TTGTATGGA CCTTTGGTCA	18240
AACCGTGGAA AATCAACCAA AAAAGTAAA GATATGGTTC GTGGGCACCA AATGGCAAAT	18300
ATGGCAGGTG TTCGTAAGCT GCAACCAAAC TTGCGTGTTC AGCCAATGGT TATCGATCCA	18360
TTTGTATCA ACGAACATAGA CTATTAACCTA GTTTCACACT TCCACAGTGA TCATATCGAC	18420
CCATACACAG CTGCAGCAAT TCTCAATAAT CCTAAGTTAG AGCATGTTAA GTTGG	18475

(2) INFORMATION FOR SEQ ID NO: 39:

- (i) SEQUENCE CHARACTERISTICS:
- (A) LENGTH: 7186 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 39:

CCAGGATTTG GTACCGTTGC AAGTGGTGTG CCTTTCTCC TAAAGGAAAA TGGAGGAAAA	60
ATCAATCAAT CAGCACATTC AGATATCAA GTTGCTAAGG TATTGGTCAA GGATGAAGAT	120
GAAAAAAATC GCTTGCTTGC AGCAGGGAAT GACTTTAAGT TTGTAACCAA TGTGGATGAT	180
ATTTTATCG ACCAGGATAT TACTATCGTA GTGGAATTGA TGGGGCGTAT TGAGCCTGCT	240
AAAACCTTTA TCACTCGTGC CTGGAAAGCT GGAAAACACG TTGTTACTGC TAACAAGGAC	300
CTTTTAGCTG TCCATGGCGC AGAATTGCTA GAAATCGCTC AAGCTAACAA GGTAGCAGT	360
TACTACGAAG CAGCAGTTGC TGTTGGGATT CCAATTCTTC GTACTTACG AAATTCTTG	420
GCTTCTGATA AAATTACGCG CGTGCTTGGG GTAGTCAACG GAACTTCCAA CTTCATGGTG	480
ACCAAGATGG TGGAAGAAGG CTGGTCTTAC GATGATGCTC TTGGCGGAAGC ACAACGTCTA	540

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GGATTTGCAG	AAAGCGATCC	GACGAATGAC	GTAGATGGGA	TTGATGCAGC	CTACAAGATG	600
GTTATTGAG	GCCAATTGCG	CTTGGCATG	AAGATTCGCT	TTGATGATGT	AGCCCACAAAG	660
GGAATCCGCA	ATATCACACC	AGAAGACGTA	GCTGTAGCTC	AAGAGCTTGG	TTACGTAGTG	720
AAATTGGTTG	GTTCTATTGA	GGAAAACCTCT	TCAGGTATTG	CTGCAGAAGT	GACTCCAACC	780
TTCCTACCTA	AAGCGCACCC	ACTTGCTAGT	GTGAATGGCG	TAATGAACGC	TGTCTTTGTA	840
GAATCTATCG	GTATTGGTGA	GTCTATGTAC	TACGGACCG	GTGCGGGTCA	AAAACCAACT	900
GCAACAAGTG	TTGTAGCTGA	TATTGTCCGT	ATCGTTCGTC	TTTGAATGA	TGGTACTATT	960
GGCAAAGACT	TCAACGAATA	TAGCCGTGAC	TTGGTCTTGG	CAAATCCTGA	AGATGTCAAA	1020
GCAAACACT	ATTTCTCAAT	CTTGGCTCTA	GACTCAAAG	GTCAGGTCTT	GAAGTTGGCT	1080
GAAATCTTCA	ATGCTCAAGA	TATTTCCCTT	AAGCAAATCC	TTCAAGATGG	CAAAGAGGGT	1140
GACAAGGGCC	GTGTCGTTAT	CATCACACAC	AAGATTAATA	AAGCCCAGCT	TGAAAATGTC	1200
TCAGCTGAAT	TGAAGAAGGT	TTCAGAAATTC	GACCTCTTG	ATACCTTCAA	GGTGCTAGGA	1260
GAATAAGATG	AAGATTATTG	TACCTGCAAC	CAGTGCCAAT	ATCGGGCCAG	TTTTGACTC	1320
GGTCGGTGT	GCTGTAACCA	AGTATCTTCA	AATTGAGGTC	TGCGAAGAAC	GAGATGAGTG	1380
GCTGATTGAA	CACCAGATTG	GCAAATGGAT	TCCACATGAC	GAGCGTAATC	TCTTGCTCAA	1440
AATCGCTTTG	CAAATTGTC	CAGACTTGCA	ACCAAGACG	TTGAAAATGA	CCAGTGATGT	1500
CCCTTGGCG	CGCGGTTTGG	GTTCTTCCAG	CTCGGTTATC	GTTGCTGGGA	TTGAACCTACC	1560
CAACCAACTG	GGTCAACTCA	ACTTATCAGA	CCATGAAAAA	TTGCAGTTAG	CGACCAAGAT	1620
TGAAGGGCAT	CCTGACAATG	TGGCTCCAGC	CATTTATGGT	AATCTCGTTA	TTGCAAGTT	1680
TGTTGAAGGG	CAAGTCTCTG	CTATCGTAGC	AGACTTTCCA	GAGTGTGATT	TTCTAGCTTA	1740
CATTCCAAAC	TATGAATTAC	GTACTCGCGA	CAGCCGTAGT	GTCTTGCTTA	AAAAATTGTC	1800
TTATAAGGAA	GCTGTTGCTG	CAAGTTCTAT	CGCCAATGTA	CGGGTTGCTG	CCTTGTGPGC	1860
AGGAGACATG	GTGACCGCTG	GGCAAGCAAT	CGAGGGAGAC	CTCTTCCATG	AGCGCTATCG	1920
TCAGGACTTG	GTAAGAGAAAT	TTGCGATGAT	TAAGCAAGTG	ACCAAAGAAA	ATGGGGCCTA	1980
TGCAACCTAC	CTTCTGGTG	CTGGGCCGAC	AGTTATGGTT	CTGGCTTCTC	ATGACAAGAT	2040
GCCAACAATT	AAGGCAGAAAT	TGGAAAAGCA	ACCTTTCAA	GGAAAACGTC	ATGACTTTGAG	2100
AGTTGATACC	CAAGGTGTCC	GTGTAGAACG	AAAATAAGA	ATAGAAGATA	GGATGGGGAA	2160
ACTCTTGACC	AGAGGGGTTTC	ATATCCTTTT	TGTGAAAAGA	AGTTTACACT	CAATGAAAAT	2220
CAAAGAGCAA	ACTAGGAAGC	TAGCCGCAGG	CTGCTAAAA	CAGTGTGTTG	AGGTTGCAGA	2280
TAGAACTGAC	GAAGTCAGCT	CAAGACACTG	TTTGAGGTT	GCAGATAGAA	CTGACGAAGT	2340

CAGTAACCAT ACTACGGTAA GGTGACGCTG ACCTGGTTTG AAGAGATTTT CGAAGAGTAT	2400
TAGTTAAAAA CGTGATAAAG GAGAAATAAA GATGGCAGAA ATTTATCTAG CAGGTGGTTG	2460
TTTTGGGGC CTAGAGGAAT ATTTTCACG CATTCTGGA GTGCTAGAAA CCAGTGGTGG	2520
CTACGCTAAT GGTCAAGTCG AAACGACCAA TTACCAAGTGTG CTCAGGAAA CAGACCATGC	2580
AGAACCGGTC CAAGTGATTT AGCAGTGAGAA GGAAGTGTCA CTCAGAGAGA TTTTACTTTA	2640
TTATTTCCGA GTTATCGATC CTCTATCTAT CAATCAACAA GGGAAATGACC GTGGTCGCCA	2700
ATATCGAACT GGGATTATTATC ATCAGGATGA AGCAGATTTG CCAGCTATCT ACACAGTGGT	2760
GCAGGAGCAG GAACGCATGC TGCGTCGAAA GATTGCAGTA GAAGTGGAGC AATTACGCCA	2820
CTACATTCTG GCTGAAGACT ACCACCAAGA CTATCTCAGG AAGAATCCTT CAGGTTACTG	2880
TCATATCGAT GTGACCGATG CTGATAAGCC ATTGATTGAT GCAGCAAACCT ATGAAAAGCC	2940
TAGTCAAGAG GTGTTGAAGG CCAGTCTATC TGAAGAGTCT TATCGTGTCA CACAAGAACG	3000
TGCTACAGAG GCTCCATTAA CCAATGCCTA TGACCAAACC TTTGAAGAGG GGATTTATGT	3060
AGATATTACG ACAGGTGAGC CACTCTTTTG TGCCAAGGAT AAGTTTGCTT CAGGTTGTGG	3120
TTGCCAAGT TTTAGCCGTC CGATTTCCAA AGAGTTGATT CATTATTACA AGGATCTGAG	3180
CCATGGAATG GAGCGAATTG AAGTTCGTT TCAGTCAGGC AGTGCCTACT TGGGTCATGT	3240
TTTCACAGAT GGACCGCGGG AGTTAGGCAGG CCTCCGTTAC TGTATCAATT CTGCTTCTTT	3300
ACGCTTGTG GCCAAGGATG AGATGGAAA AGCAGGATAT GGCTATCTAT TGCCCTACTT	3360
AAACAAATAA AACAGAGAGT GGGGCTTCCC ACTTTCTTCA TTTCTAGAAT ATGAATAGAA	3420
GGGATTTATG AAACACCTAT TATCTTACTT CAAACCCCTAC ATCAAGGAAT CAATTTAGC	3480
CCCCTTGTTTC AAGCTGTTAG AAGCTGTTTG TGAGCTCTTG GTTCCCATGG TGATTGCTGG	3540
GATTGTTGAC CAATCTTAC CTCAGGGAGA TCAAGGTCTAT CTCTGGATGC AGATTGGCCT	3600
GCTCCTTATC TTTGCAGTAA TTGGCCTTTT AGTGGCCTTG ATAGCTCAAT TTTACTCAGC	3660
AAAGGCAGCA GTAGGTTCTG CTAAGGAATT GACAAACGAT CTTTATCGTC ATATTCTTC	3720
CTTGCCCAAG GACACCAGAG ACCGTCTGAC AACTTCTAGT TTGGTCACTC GCTTGACTTC	3780
GGATACCTAC CAGATTCAAGA CTGGTATCAA TCAATTCCCTG CGTCTCTTTT TACGAGCGCC	3840
CATTATCGTT TTTGGTGCCA TTTTTATGGC TTATCGAATC TCAGCTGAGT TGACTTTCTG	3900
GTTCTTAGTC TTGGTTGCCA TTTTGACCAT TGTCATTGTA GGGTTATCTC GATTGGTCAA	3960
TCCTTTCTAC AGTAGTCTCA GAAAGAAAAC GGACCAACTG GTTCAGGAAA CGCGCCAGCA	4020
ATTGCAAGGG ATGCCGGTTA TTCGTGCTTT TGGTCAAGAA AAACGAGAGT TACAGATTTT	4080

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TCAAACCCTT AACCAAGTTT ATGCTAGATT ACAAGAAAAG ACAGGTTCT GGTCTAGTTT	4140
ATTAACACCT CTGACCTATC TGATTGTCAA TGGAACTCTT CTCGTTATTA TCTGGCAAGG	4200
CTATATTTCA ATTCAAGGAG GAGTGCTAG TCAAGGTGCT CTCATTGCTC TTATCAATT	4260
CCTCTTACAG ATTTGGTGG AATTGGTCAA GCTAGGCCATG TTGATCAATT CCCTCAACCA	4320
GTCCTATATC TCAGTCAAGC GAATCGAGGA AGTCTTGTT GAGGCTCCAG AGGATATCCA	4380
TTCAGAGTTA GAACAAAAGC AAGCTACCAG AGATAAGGTT TTACAAGTCC AAGAATTGAC	4440
CTTTACCTAT CCTGATGCGG CCCAGCCTTC TCTGAGATAC ATTTCCCTTG ATATGACTCA	4500
AGGACAAATT CTAGGTATCA TCGGGGAAAC TGGTTCTGGT AAATCAAGCT TGGTGCAACT	4560
CTTACTTGGA CTTTATCCAG TAGACAAGGG GAACATTGAC CTTTATCAAATGGACGTAG	4620
TCCTCTTAAT TTGGAGCAGT GGCGGTCTTG GATTGCCTAT GTACCTCAAA AGGTCGAAC	4680
CTTTAAAGGA ACCATTCGTT CCAACTTGAC TCTAGGTTTC AATCAAGAAG TATCTGACCA	4740
GGAACTCTGG CAGGGCTTGG AGATTGCGCA AGCTAAGGAT TTTGTCAGTG AAAAGGAAGG	4800
ACTCTTGGAT GCTCTAGTTG AGGCAGGGGG GCGAAATTTC TCAGGTGGAC AAAAACAAAG	4860
ATTGTCATC GCCCGAGCAG TCTTGCCTCA GGCTCCGTTT CTCATCCTAG ATGATGCAAC	4920
CTCGGCCTG GATACCATTA CAGAGTCAA GCTCTTGAAA GCTATTAGAG AAAATTTC	4980
AAACACGAGC TTAATTTGA TCTCTCAACG AACCTCAACT TTACAGATGG CGGACCAGAT	5040
TCTCCTCTTG GAAAAAGGTG AGTTGCTAGC TGTTGGCAAG CACGATGACT TGATGAAATC	5100
CAGCCAAGTC TATTGTGAAA TCAATGCATC CCAACATGGA AAGGAGGACT AGAATGAAAC	5160
GACAAACTGT AAACCAAGACG CTCAAACGTT TAGCCGTAGA TTTAGCAAGC CATCCTTTCC	5220
TCCTTTCTCT AGCCTTCTA GGAACTATTG CCCAAGTGG CTTATCAATT TACCTACCTA	5280
TTCTGATTGG GCAGGTCATT GACCAAGTCC TAGTGGCTGG TTCATCACCA GTTTTTGGC	5340
AGATTTTCTT CCAGATGCTC TTGGTGGTAA TAGGAAATAC TCTGGTACAA TGGGCAATC	5400
CTCTCCTCTA TAATCGTCTA ATCTTCTCTT ATACCAAGAGA TTTACGGGAG CGAATCATCC	5460
ATAAGCTCCA TCGTTTACCG ATTGCTTTTG TAGATAGGCA AGGTAGTGGA GAGATGGTTA	5520
GTCGTGTAAC CACGGACATC GAACAGTTGG CAGCTGGCTT GACCAGTATT TTTAACCAAT	5580
TTTCATTGG TGTTTGATG ATTTTGGTCA GTATTCTAGC CATGCTCAA ATTCACTCC	5640
TCATGACTCT CTTAGTCTTG CTGTTGACGC CACTGTCAT GGTGATTCA CGCTTTATTG	5700
CCAAGAAATC CTATCATCTC TTCCAGAAGC AAACAGAGAC GAGGGGAATT CAGACTCAGT	5760
TGATTGAAGA ATCGCTTAGT CAGCAGACTA TAATCCAGTC CTTCAATGCT CAAACAGAAAT	5820
TTATCCAAG ATTGCGTGAG GCTCATGACA ACTACTCAGG CTATTCTCAG TCAGCCATCT	5880

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TTTATTCTTC AACGGTCAAT CCTTCGACTC GCTTTGTAAA TGCACTCATT TATGCCCTTT	5940
TAGCTGGAGT AGGAGCTTAT CGTATCATGA TGGGTTCAGC CTTGACCGTC GGTCGTTAG	6000
TGACTTTTTT GAACTATGTT CAGCAATACA CCAAGCCCTT TAACGATATT TCTTCAGTGC	6060
TAGCTGAGTT GCAAAGTGCT CTGGCTTGCG TAGAGCGTAT CTATGGAGTC TTAGATAGCC	6120
CTGAAGTGGC TGAAACAGGT AAGGAAGTCT TGACGACCAG TGACCAAGTT AAGGGAGCTA	6180
TTTCCTTAA ACATGCTCT TTTGGCTACC ATCCTGAAAA AATTTTGATT AAGGACTTGT	6240
CTATCGATAT TCCAGCTGGT AGTAAGGTAG CCATCGTTGG TCCGACAGGT GCTGGAAAAT	6300
CAACTCTTAT CAATTCCTT ATGCGTTTT ATCCCATTAG CTCGGGAGAT ATCTTGCTGG	6360
ATGGGAATC CATTATGAT TATACACGAG TATCATTGAG ACAGCAGTTT GGTATGGTGC	6420
TTCAAGAAC CTGGCTCACA CAAGGGACCA TTCATGATAA TATTGCTTT GGCAATCCTG	6480
AAGCCAGTCG AGAGCAAGTA ATTGCTGCTG CCAAAGCAGC TAATGCAGAC TTTTCATCC	6540
AACAGTTGCC ACAGGGATAC GATACCAAGT TGGAAAATGC TGGAGAAATCT CTCTCTGTCG	6600
GCCAAGCTCA GCTCTTGACC ATAGCCCCAG TCTTTCTGGC TATTCCAAG ATTCTTATCT	6660
TAGACGAGGC AACTTCTTCC ATTGATACAC GGACAGAAAGT GCTGGTACAG GATGCCTTTG	6720
CAAAACTCAT GAAGGGCCGC ACAAGTTCA TCATTGCTCA CCGTTGTCA ACCATTCAGG	6780
ATGCGGATTT AATTCTTGTC TTAGTAGATG GTGATATTGT TGAATATGGT AACCATCAAG	6840
AACTCATGGA TAGAAAGGGT AAGTATTACC AAATGAAAAA AGCTGCGGCT TTTAGTTCTG	6900
AATAAGCCAT TCTCTTTGAA AGTTTATGG ACGAAAAAAAG TTGCTTCGA GTGACTTTTT	6960
TGTTACAATA GCTAGAAAAA TTGTTCACTG TAATACTCAA TGAAAATCAA AGAGCAAAC	7020
AGGAAGCTAG CCGTAGGTTG CTCAAAGCAC AGCTTTGAGG TTGTAGATAA GACTGACGAA	7080
GTCAGTTCAA AACACTGTT TGAGGTTGCA GATAGAACTG ACGAAAGTCAG CTCAAAACAC	7140
TGTTTGAGG TTGCAGATAG AACTGACGAA GTCAGCTCAA AACAGG	7186

(2) INFORMATION FOR SEQ ID NO: 40:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 14273 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 40:

CTGAAAATTC TAAAAAATTT ATAAGTAAGG AATTAATTAG TTATTTTGT GATAAAAGTTT	60
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ATGATGAAAT ATTTGTTGAA GAGGTAGTTC CGCACGTTT TCTGCCATAT GAATCTGACT	120
TACTTCTTAT TTTACCGACT ACGGCAAATG TGATTGGCAA AATTGCTAAT GGTATTGCTG	180
ATGATTAGT TACAGCAACT GTTTTAAACT TTAAATAAAA AATAATTTT TGTCCTAATA	240
TGAACCTCAC TATGTGGGC AACACATAG TTCAAAGAAA TGTATCAATT CTAAGGAGT	300
TGGGACATAT ATTTTATTT GAGTCTAAAA AACATATGA GGAGGATTG CGTAAAGCAA	360
TAGATTCAAC ATGTCATG TTACAACCAC AGTCGTTAGT AAAAGAACCT ATCAAATTAG	420
AAAATATTGT CCTTGAAGAG GGACATTTAA AACTACTGAG AATATTAATG AGGGGAAAAAA	480
ATGGAAAATT CATCAATCGA TGTAGATATG CTGTTGGAAAG AATTGACACA AGAAGCAATG	540
GTCGTTGTTG CTGTTGATAA GGACTGTTAA TTTAAACTTA TGGCAATATA TGAAAGGTTA	600
CTGGATGTTT TAAATTATGC AGGAGTAGC CTTTATTAT ATACAAATGG ATAAAGTAAG	660
GATAATACAA TGATTAATAA AAAATACAA CAAGTTGTT TGGAATCATT ACAGAATT	720
TTGAATGGGA ACTTCATTT GCCTTGTTA GTCTATGATT TTGGCTTGCT GGAAACTGTA	780
CTTGATGAAT TTAAAAATCA AATTCTGTA ACATTCAATT ACCAACTTTT TTATGCCGTT	840
AAAGCAAATT CAAATGAGAA GATACTTGAA TTCTTAGTAG ATAAAATTGA TGGAGTTGAT	900
GTGGCGTCAT TATCTGAATT AGATGTGGCT AAAAATTGTT TCCCACCAAC TCAAATTCT	960
GTTAATGGTC CCGCATTTTC TTATGAAACT TTATATAATC TGATTAAAA ACAATATAAA	1020
GTTGATATTA ACTTTTGGA ACATCTCAA CAAATTTCC CAAAGAAC TGTGGAATA	1080
AGAGTAACGG AGCCAGATGAA ACTTAATAAT CGTATGAGTC GATTGGAAT AAATATTG	1140
ACTGATAATT GGACTAGTAA TTTACAAAT CCTTTAATTA CACGACTGCA TTTCAATT	1200
GGAGAAAAG ATGATAATT TATTGTTAAG TTAGATAAAA TATTATTTAA GTTACAAGAA	1260
ATTAATAAAC TTAGAGAGGT TAGAGAAATA AATCTGGAG GCGGTTTTAT GAAATTATT	1320
ATGGAAAATC GTTGAAAGA ATTTCCTCTA TCACCTATGG AAATCTATAA AAAGTACGAT	1380
ATTGATAGTA CTGTGACTAC AATAATAGAA CCAGGTAGTG CAATTACTTC ATTTCTGCC	1440
TATATGATTA CTAGCCCCAGT TAATGTTAGT GAGGTGAATG AGCAGCAGGT TATCACGTTA	1500
GACACATCAA TATACACCAA TACATTATGG TTTGTTCCGC ATATTATTAC AACGTTAAAT	1560
TCAAGTAGTA AAGAGCGTTA TAGTACTATT CTCTATGTTA ATACCTGTTA TGAACATGAC	1620
AACTATAAAA TGAAAGTTTC GCTTCCAAGG TTAACCTCAA ATAGCAGTAT AGTGT	1680
CCTGTAGGAG CTTATATAAA AACCAATCAT TCAAATTAC ATCGTAATGA TTTTATGCCG	1740
GAGGTATATT TGTGGACAAA AAACCTTGACA TATTAGATAA AGTTAAGGAA TATTTAGGAA	1800
ATAAAACCTAC TCAAATTCTG GATAATCAAT ATAAAGAATT TTTGAAACTT AATGATATAA	1860

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GGCGAGCGTT TGGTATTC AAAAAAGTAT TAAACAATTC TTTTAATTTC ACGAGTAAAG	1920
AATTTAATGA TTTAATTAAT AACGAAAATT ATTTATTGCA ATATGCATGT AGAATTAGAG	1980
AGGAATGGAG AAAAAAAATGC TTTAACATT CTTATCGTT TCTATGCTCA CCTATAATTA	2040
CAGATGATT TCTAACACG AAGACATTGA GAAGTAGCCA AATTGAATAT AAATATGAGC	2100
GATATTTATC GAAAAGTCG ATAGGCGATA GAGCGGTTGA TGGCTTTGTT TCCTTCAATA	2160
CTTTAACAGC TAATGGTATG TCTGCTATTA AACTATGTCT TGAGATATTA AACTCTATTT	2220
TCTTCAAGAA GAAGATTGAT TTATTATATT CAACCGGATA TTATGAAACA AGATTTTAT	2280
TAAATAATCT TGCTAAATCA GGTATTAGTT GCTATGAGGT AAGTAATTGT GAATTGGATA	2340
AAGATAAAATT TTATAATGTA TTCATGATGG AACCAATCG AGCCGATTAA ACATTACAAA	2400
AAACTGATT CAAGATAGTA CAATATTTG TTAAGTATAA AAATAATTCA ATAAAAGTCG	2460
TTATTTAGA TATTCATAT CAAGGTTCTA ATTTAAATT AGTAGAATT TTAGAGAAAT	2520
TTAAATTGTC GAATGTAATT ATTMTTGTTG TAGCATCTT GATAAAATTA GATCAAATGG	2580
GATTAGAATT GACAAATGGG GGAATAATAG AAGTGTTTAT TCCTAATCAT TTGAGAAAGT	2640
TGAAAAAATT TATTGAAGAG GAATTCAATA AATTAGAAA TTCTCACGGA GCTAATCTAA	2700
GCCTCTATGA ATACTGTTG CTTGATAATT CTTAACCTT AAAAAATGAT TGGAACATT	2760
CTGATTAGT TATGAAATT ACCAGTAATT TTTATGCTGA TATAAAAGAC TTGTTCATGG	2820
AAAATTCTGA TATTGAAATC ATCCATGAAG AGGGAGTACC TTTTGTATTT TTAGATTTAA	2880
TAGGTGAAGG TAAAAAGAA TATGAAATGT TTTTCAATG GTTAAACTTC TTTTACAAAC	2940
AGCTTGGAAAT CACATTGTAT CCTAGAAATA GTTTGGGTT TCGGAATCTA ACAGTAGAGT	3000
ATTTTGGAAAT TATTGGGACA GAAAGATATA TATTTAAGAT TTGTCCAGGT GTTATCAAAG	3060
GGTTAAGTTA TTATTTGATG AAATTTTTAT TAAATCTTT TTCAAATGAA TATTTAAAAA	3120
CTACTGATGA GGTTAATAGA TGAAAAATTG GATAAAGTTG CTAATAATTAA GATTGATTGT	3180
TAACTTAGCA GACAGTGTAT TTTATATAGT ACCATTGTTG CACGTTAGCA ATAATTATTC	3240
TTCGACCATG TTCTTAGGAA TATTTATTGCA ACTAAATTAT CTACCGGATT TGTTACTAAT	3300
CTTTTTGGAA CCAGTTATG ACAGAGTAA TCCGAAAAA ATTCTTATAA TATCAATT	3360
GGTTCAATTAA GCAGTGGCTG TAATATTTT ATTATTATTA AACCAAATAT CATTGGGT	3420
GATAATGAGT CTAGTGTAA TTTCAGTAAT GGCTAGCTCC ATAAGTTACG TGATAGAAGA	3480
TGTGTTGATT CCTCAAGTGG TAGAATATGA TAAGATGTAA TTTGCAAATT CTCTTTTAG	3540
TATTCGATTA AAAGTATTAG ATTCTATTTT TAATTCAATT GCATCATTTC TACAGGTGGC	3600

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AGTAGGATTT ATTTTATTGG TTAAGATAGA TATAGGCATA TTTTTACTTG CTCTATTTAT	3660
ATTGTTGTTG TAAAGATTTA GAACTAGCAA TGCGAATATA GAAAACCTCT CTTTCAAATA	3720
TTACAAGAGA GAACTGTTGC AAGGTACAAA GTTTATTTA AATAATAAAAT TATTATTTAA	3780
AACCACTATT TCTTTAACGC TTATAAACCTT TTTTTATTCA TTTCAGACAG TAGTTGTACC	3840
GATTTTTCT ATTGATATT TTGATGGTCC GATTTTTAT GGTATTTTT TAACATTGC	3900
TGGTTTGGGT GGTATATTGG GAAATATGCT AGCGCCAATC GTAATAAAAT ATTTAAAATC	3960
GAATCAAATT GTGGGTGTAT TTCTTTTTT GAACGGCTCA AGTTGGTTAG TAGCAATTGT	4020
TATAAAAGAC TATACTTTAT CACTTATTGT ATTTCGTT TGTTTTATGT CTAAAGGAGT	4080
CTTCAATATT ATTTTAATT CGTTGTACCA ACAAAACCT CCACATCAAC TTCTGGTAG	4140
GGTAAATACT ACCATTGATT CTATTATTC TTTTGGATG CCAATTGGTA GTTTAGTTGC	4200
AGGAACGCTT ATTGATTGTA ATATTGATT AGTGTAAATT GCTATTAGCA TACCTTATT	4260
TTTGTCTCT TATTTTTTAT ATACGGATAA TGGATTGAAA GAATTTAGTA TATATTAGAA	4320
ATGTTATGT TCATTCAAAA GCATAATGAC TATAACTGAA AAAGAAAAGT GATATCTTA	4380
AGGTTGTTCT TCTTGGTGGT GAGATTGCGT AGACAAACCA AGCTTTGTC GGAAAGATTA	4440
CCAATGCTTT GATGGATAGG ATGTAATTGTA GCAAGATGTT TTTAGTGGTA ACGGTATCGT	4500
GGATGGACGT GAAATAACCT CTTCTTCGA GGAGTATTT ACTAAAAAAC TAGCCTPGGA	4560
GCGTCCCCA GAAACGGACT TACTCATTTGA CTCTTCAAAC ATTTGGGAG AAGATTTGC	4620
TTCATCTGTT CCTTGAAAAA AGTCACAGCA GTCATCACAG ACGATAGTAC TGAACAAAAC	4680
TATGAAGAGT TAGAAATTAA TACGCAGGTG ATTGTATAA GGATCTGGAA ATAGATAAGA	4740
AGTTGATTAG TATTGACCTA CGTGGTACAA ATATTAAGAT TACTGTTCTT TCAAATGACG	4800
GTGAGATTGA AACCTTGTGG AGTATTACAA CAGATACAAG TGAGAAAGGT TCTCAAATT	4860
TATGGACAT CATCAGTTCT ATTAACAAATA AATTGACCGA ACGGAATATT CCTGATAGCG	4920
ACCTCTTGG AATCGGTATG GGAAGTTGCT CATCATACTT TCCTTGAAA TCATAGGGC	4980
TATAAACTCT CCGTCTACTT GTCCTGCAAC AATTGAAGTC TGCTAAACCGCT	5040
AATCTTTCA TAGACTTTCT CCCTTTAGG AGCCTAGCTT TCTAGTTGT TCTTGATTT	5100
TTATTGAGTA TACCACTATT TTACTCCCTC TGGCAAGGGA CTTTGTCTAT CTGGAGGGAT	5160
TGGGCTCTTA TGTGGTGGAG CTTTCTGTT CTTTCTGAA TATGGTATAA TAGCACTAAT	5220
CAATTCTAG GAAAATAGAT ACAGAAAGGG GCTGAAAGAT GTCTCATATT ATTGAATTGC	5280
CAGAGATGCT GCCAAACCAA ATCGCGGCTG GAGAGGTCA TGAACGTCTT GCCAGTGTGG	5340
TCAAAGAGTT GGTAGAAAAT GCCATTGACG CGGGCTCTAG TCAGATTATC ATTGAGATTG	5400

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AGGAAGCTGG TCTCAAGAAG GTTCAAATCA CGGATAACGG TCATGGAATT GCCCCACGATG	5460
AGGTGGAGTT GGCCCTGCCT CGCCCATGCAGA CCAGTAAGAT AAAAAATCAA GCAGATCTCT	5520
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AGGATCTCTT TTTCAACACG CCTGCCCGTC TCAAGTATAT GAAGAGCCAG CAAGCGGAGT	5760
TGTCTCATAT CATTGATATT GTCAACCGTC TGGGCTTGCC CCATCCTGAG ATTTCTTTA	5820
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ACCTAGATTT CGAAATTTCAG GGGTTTGTGT CCTTGCCCTGA GTTGACTCGG GCTAACCGCA	6000
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AACAAACCTT GATTCCAGAT GCCTTGAAA ATCTTGCCAA ATCGACCGTG CGCAATCGTG	6300
AGAAGGTGGA GCAAACCTATT CTCCCACCTCA AAGAAAATAC GCTCTACTAT GAGAAAACCTG	6360
AGCCGTCAAG ACCTAGTCAA ACTGAAGTAG CTGATTATCA GGTAGAATTG ACTGATGAAG	6420
GGCAGGATTT GACCCCTGTTT GCCAAGGAAA CCTTGGACCG ATTGACCAAG CCAGCAAAAC	6480
TGCATTTGC AGAGAGAAAG CCTGCTAACT ACGACCAGCT AGACCATCCA GAGTTAGATC	6540
TTGCTAGCAT CGATAAGGCT TATGACAAAC TGGAGCGAGA AGAAGCATCC AGCTTCCCAG	6600
AGTTGGAGTT TTTCGGACAA ATGCACGGGA CTTATCTCTT TGCCCAAGGG CGAGATGGAC	6660
TTTACATCAT AGATCAGCAC GCTGCTCAGG AACGGGTCAA GTACGAGGAG TACCGTGAAA	6720
GCATTGGCAA TGTTGACCAA AGCCAGCAGC AACTCCTAGT GCCCTATATC TTTGAATTTC	6780
CTGCGGATGA TGCCCTGCCT CTCAGGAAA GAATGCCCTCT CTTAGAGGAA GTGGGCGTCT	6840
TTCTAGCAGA GTACGGAGAA AATCAATTAA TTCTACGTGA ACATCCTATT TGGATGGCAG	6900
AAGAAGAGAT TGAATCAGGC ATCTATGAGA TGTGCGACAT GCTCCCTTTG ACCAAGGAAG	6960
TTTCTATCAA GAAATACCGA GCAGAGCTGG CTATCATGAT GTCTTGCAAG CGATCTATCA	7020
AGGCCAATCA TCGTATTGAT GATCATTCAG CTAGACAACT CCTCTATCAG CTTTCTCAAT	7080
GTGACAATCC CTATAACTGT CCTCACGGAC GTCCCTGTTT GGTGCATTTC ACCAAGTCGG	7140

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ATATGGAAAA GATGTTCCGA CGTATTCAAG AAAATCACAC CAGTCTCCGT GAGTTGGGGA	7200
AATATTAAAA GTATAAAAAA GTCTGGAAA AATTTTCAAA ATCAAAAAAA CGCATAAAAT	7260
CAGGTGTTCA AAAACCTTGA TTTTATGCCGT TTTATCATGG AAATAGTTAC TTCATTTTT	7320
CCTAATTCTT TTGCAAACTC TTTTAAACG ACGTCAGTTT TATCAGTAAT CTCAAAACAG	7380
TGTTTGAGC TAATTTGCC AGTTTTGTCT GTAACATCGA AGTTGTGTT TACCACTCTG	7440
CGACTGGTT CCTAGTTGC TCTATGATT TCACAGAGCA TTAAATTGCG ATTTTGCCAA	7500
GTTTCTTAT TCGTCTAAA GTAGAGTCTG TTCTATGCCGT CTAATGTACG AATCAGGTTG	7560
ACCATTCAA TAGCTCCTTG TGCAACTCA GAACCCATTAT TTCCCTGCTTT AGTACCAAGCT	7620
CGTTCTATGG CTTGTTCAAT TGTATCTGTC GTTACACAC CAAACATAAC AGGAATTTCG	7680
CTATTAAAC TGATTTGGGC GATTCCCTTA GATACCTCGC TACATACATA ATCATAATGA	7740
CTTGTATTCC CTCTAATGAC AGCTCCCAAG CAGATAATTG CATCATATT TTTACTTTT	7800
GCCATTTTG ATGCAATCAG TGTTTTC AAAGCTCCTG GAACCCAGGC TACCTCTATA	7860
TCTTTCTCGT TTACATTCTC TCTTTGAGA TTATCTAGTG CTCCAGATAA TAATTTGAA	7920
GTTATAAATT CATTAAATCT CGCTACAACA ATACCTATTT TAATATTGTT TGCTACTAAA	7980
TTACCTTCAT AAGTGTTCAT TTATTTTCC TCCATATTAA AAATGTGACC CATTGATT	8040
TTCTTGTTT CTAAATAAAA ACTATCGAA GGATTTGGCTT CTATTCGAT TGATATTCTA	8100
CTGGAAATGG TAATTCATAA TTGTTCTAAC TGTTCAACCT TGTCACCAATT ATTTGTCAGT	8160
AAATGAAGTG ACTGAAGTCC CAGATCTTA AGCATTGTTG CTCCAATATG ATATTCTCTT	8220
AAATCACCTT CAAAGCCTAA TGCAAGATTG GCATCAAGCG TATCCATGCC TTGATCTTGT	8280
AAATGATAGG CTTTTAATTG ATTGATAAGT CCAATTCCCTC GTCCCTCCTG TCGCAAGTAA	8340
AGTAAGACAC CCGAACCAATT CTCAACAAATC ATTTTCATAG CCTTATCGAA TTGCTGTCCA	8400
CAATCGCAAC GTAAAGAGCC TAAAACATCT CCTGTTAAC ATTCCGGAGTG GACCCGACAT	8460
AATACATTGG CTTCATCCCTC TATATTTCCC ATAATAAGAG CAAGATGATG TTCCCCATT	8520
AGTTTATCTA TATAGCTAAT TGCTTGAAA TTACCGTATC TAGTAGGCAT ATTGACAGTT	8580
GAAACTCGTT CTACCAAGCTG ATCATATACT TTTCTATATT CTTGTAATTG TTTGATGGTA	8640
ATTAGTGGAA TGTTGTGTTT TTTCGAGAAC TGAATTAAAT CATCTGTTCT CATCATTTG	8700
CCATCATGAT TCATTATTTCA ACAACATAGG CCACACTCTT TTAGTCCAGC TAATTTAAT	8760
AAATCAACAG TTGCTCTGT GTGTCCTATT CTTTCTAGGA CACCACCTTT TTTGCAATT	8820
AAAGGAAACA TGTGTCCTGG CCTGCGAAAA TCAGAGGGTG TTATATCTTC AGCTACACAC	8880
ATACGTGCGG TCAGTCCTCT TTCCCTGGCA GAAATACCTG TGTCGTTTC TTTATAATCA	8940

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ATTGAAACTG TAAAAGCAGT CTTATGATTA TCTGTATTGT TTTCAACCAT AGGTGAAAGC	9000
ATTAATTGAT TAGCTAAACT TTCGCTCAT A GGCAACAAA TTAATCCTT GGCATAAGTA	9060
GCCATAAAAT TAACATTTTC TGTTGTAGCT GCTTGTCAG AACAAATTAA GTCTCCTTCA	9120
TTTTCTCTAT CCTTGTGTC TATAACAAGA ACAAGTCGTC CCTTCTGCAA TGCTTCTAAT	9180
GCTTCTTGT A TTTTCGATA TTCCATTGAC TGATTATCCT TTCTGCTAA ATCCATTTG	9240
ATATAATAGT TCCTTAGATA TTTCTGATTT TGGAGAGTTA TCCATCAGTT TPTGCACATA	9300
TTTACCTAAG ATATCATTTC CAAGATTAC TGTACTCCCG ACTTGTTAC TCTTAAGAAT	9360
GGTTMTTCC AAGGTATGAG GGATAACAGA TACTGAAAG TTTACTTTGG AGACTTTAGC	9420
GACAGTCAGA CTAATGCCGT CAATTGTAAT AGATCCTTT TCAACTATTA AATCTAAAAT	9480
TTCTTTTTGT GTGTTGATTT GATACCATAAC AGCATTATCA TCTTTTTTA TTGACGAGAT	9540
TTTTCTGT A CCATCAATGT GTCCTGTAAC GACGTGACCC CCAAGTCGAC CGTTGACAGA	9600
TAAGGCTCTT TCTAGATTCA CCTCACTTCC ATGTTTTAAT AGAGTAAGAG CTGTTGACT	9660
CCATGTTCA TTCATTACAT CAACTGTAAGG GGATTGATGA TTGAAATGAG TAACTGTAAG	9720
ACAGATACCA TTTACTGCTA TACTATGCC TAAATGGATA TCCGTTAATA TTTTGAGGC	9780
TTTAATTGAT AGTTTACAAT TACGAGAGTC TTCTGTATT CTTTCAACTT TTCCGATTTC	9840
TTCAATTATT CCTGTGAACA TGGATAAAC ACTTCACTTT CTATGAGATA GTCATTTCT	9900
ATTTGAGAAA ATGCATAAGG TTCAATCTA ATAGCGTCAT TTGGCAAAGA AATACCTTCA	9960
CCTCCGACAG GAAACCTGGC ACTACCTCCA AAAACCTTTG GTGCAATATA TATTTTCAGC	10020
TCATCAACAA TTTGTTGTC CAAAGCACTC CAATTCAATTA GACTGCCCTT TTCTAGAACT	10080
AGGCTATCAA TCTGCATGTT TCCTAGATGT TGCATTAAC TCGATAAGTC TATATGATTG	10140
CCTTTTTCT TTATGGAAAG TATTCACAG CCATGATTTT GATATAGCTT CATTTTATTG	10200
TTGTCCTTCAG AGGAAGTGGC AATGTAAGTT TTAATATCAT TTGCTGTTTT TACGATTTA	10260
GAGGTAAGAG GAGTCGCTAA ATGTCATCG CATATGATAC GGATAGGATT TTTCCCTTCC	10320
TCCAATCTAC ATGTCAGCAA AGGATCGTCT TGAATAACAG TATTGACTCC CACCATAATT	10380
GCACTAACAT GGTGTCGTAAC TGATGCACTA TGCTTCTTG CTTCTCTTC AGTAATCCAT	10440
TTGGATTTGAT TTGTTTTAGT GGCTATTTTT CCATCCATTG ACATTGCTA TTTCTAAAAA	10500
ACATAGGGTA CATGCTGGGT AATATACTTT CTAAAACCTT TTATTAAGTT AAGACACTCA	10560
TTTTCTAAAA TTCCAACAGT AACTTGAGA TTATTTCTT CAAGTATCTT TACTCCTTTT	10620
CCAGATACAA TAGGATTACA GTCTAGGCTT CCAATGACTA CTCTTGTAA ACCACTATCG	10680

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ATTATAGCAT CTATACAGGG AGGTGTTTC CCGAAGTGAC AACAGGGTTC AAGTGTACA	10740
TAAAGCGTCG CTCCGACAGG GGATTCTCTA CAGTTTTAA GAGCATTCT CTCAGCATGT	10800
GGGCCACCAA AAAACTCATG ATAACCTTGT CCGATAATGT GATTATCTTT TACAATAACT	10860
GCGCCGACCA TAGGATTGGG ATTGACGTAA CAGCCCCCTT TTTGTGCCAG TTTTATTGCT	10920
AATTCATAT ATTTGAATC GCTCATCTCG CTACCTCCAA AAAATATAC CTTGAATAGG	10980
GGACTACTCA AGGCATACAA AAGAAAACCTT ATGCGATTAA CAAAAATGCT CTGAAATGAC	11040
AAGTAATCAT TTCAGAGCAC GCAAAAAGCA CAAATATACT TTTATCTTCT TTCACTCCAGA	11100
CTATACTGTC GGCTTGGAA TTTCACCAAA TCATGCCTTT CGGCTCGTGG GCTATACCAC	11160
CGGTAGGGAA TTTCACCCCTG CCCTGAAGAT AGTTATTCAA TTACAGATGA TTATAGTACT	11220
TAATTTGAA TATGCAACA GATAAAATACC GATTGTTTT GATATACTGT ATTTGTGATA	11280
ATCGATTCTC GCTCCTCGGA TAAAGAAAAT ATGATATACT AGATAAACGA AATAAGAGAG	11340
AAGGAATACT ATGTACGCAT ATTTAAAAGG AATCATTACC AAAATTACTG CCAAATACAT	11400
TGTTCTTGAA ACCAATGGTA TTGGTTATAT CCTGCATGTG GCCAATCCTT ATGCCTATTC	11460
AGGTCAAGGTT AATCAGGAGG CTCAGATTAA TGTCATCAG GTTGTGCGTG AGGACGCCA	11520
TTTGCCTTAT GGATTTGCT CAGAGGATGA GAAAAGCTC TTTCTTAGTC TGATTTGGT	11580
CTCTGGGATT GGTCTGTAT CAGCTCTTGC TATTATCGCT GCTGATGACA ATGCTGGCTT	11640
GGTTCAAGCC ATTGAAACCA AGAACATCAC CTACTTGACC AAGTTCCCTA AAATTGGCAA	11700
GAAAACAGCC CAGCAGATGG TGCTGGACTT GGAAGGCAAG GTAGTAGTTG CAGGAGATGA	11760
CCTTCCTGCC AAGTCGCAG TGCAAGCAAG TGCTGAAAAC CAAGAATTGG AAGAAGCTAT	11820
GGAAGCCATG TTGGCTCTGG GCTACAAGGC AACAGAGCTC AAGAAAATCA AGAAAATTCTT	11880
TGAAGGAACG ACAGATACAG CTGAGAACTA TATCAAGTCG GCCCTTAAAA TGTTGGTCAA	11940
ATAGGAGCAG AGAATGACAA AACGTTGTTG GTGGGTCAAG ATGACCAACC CGCTCTACAT	12000
CGCCTATCAT GATGAGGAGT GGGGCCAGCC CCTCCATGAT GACCAAGTAT TGTTTGAGTT	12060
GTTGTGTATG GAAACCTATC AGGCAGGGCT GTCTTGGAA ACGGTACTCA ACAAACGCCA	12120
AGCTTCCGA GAAGTCTTTC ATAGCTATCA AATTCACTCA GTCGCAGAGA TGACTGACAC	12180
TGAATTGGAA GCCATGCTGG AGAATCCAGC TATCATTGCA AATAGAGCCA AGCTTTTGC	12240
TACACGCCGT AACGCCAAG CCTTTCTACA GTTACAGGCA GAGTACGGCT CTTTGATGC	12300
CTATCTTGG TCTTTGTTG AGGGGAAAAC TGTCGTTAAC GATGTTCCCTG ATTATGCCA	12360
AGCGCCAGCT AAAACACCCCT TATCTGAGAA ATTAGCCAAA GATCTAAAA AACGAGGCTT	12420
CAAGTTACACA GGCCCAGTCG CCGTATTGTC TTTCTACAG GCTGCAGGGC TAGTTGATGA	12480

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CCACGAGAAT GATTGTGAGT GGAAAGGTCT TAAATGATGT CTAACAAAAA TAAGGAAATT	12540
CTGATTTTG CGATTCCTCA TACAGTCCTC TTTATGTTTG ATGGCGTTAA ATTGCTGGCT	12600
TCTTTAATGC CATCTGCCAT TGCAAATTAT CTTGTTATG TAGTTTAGC TCTATATGGC	12660
TCCTCTTGT TCAAGGATAG ATTGATCCAA CAATGGAAGG AGATTAGAAA GACTAAAAGA	12720
AAATTCTCT TTGGAGCTT AACAGGATGG CTCTTCTCA TTCTGATGAC TGTTGTCTT	12780
GAATTGTAT CAGAGATGTT GAAGCAGTT GTGGGACTAG ATGGACAAGG TCTAAATCAG	12840
TCTAATATTC AAAGTACCTT TCAAGAACAA CCACTACTGA TAGCTGTTT TGCTTGTGTC	12900
ATTGGACCTC TGGTAGAAGA ATTATTTTC CGTCAGGTCT TATTGCATTA CTTGCAGGAA	12960
CGGTTGTCAG GTTTACTAAG CATTATTCTG GTAGGACTTG TTTTGCTCT GACTCATATG	13020
CACAGTTGG CTCTATCAGA GTGGATTGGT GCAGTTGGT ACCTAGGTGG AGGCCTTGCC	13080
TTTTCTATTA TTTATGTGAA AGAAAAAGAG AATATCTACT ATCCCCTACT TGTTCACATG	13140
TTAACGCAACA GCCTCTCCTT AATCATTAA GCTATCAGTA TAGAAAAATG AAATGAGAAC	13200
AGGACAAATC GATTCTAAC AATGTTTAG AAGTAGAGGT GTACTATTCT AGTTCAATA	13260
TACTGTAATA TGTGATGAAA ATGCCAGTAA TGATACCGAG AAAAAGCTG AGAAACTTT	13320
CCCAGCTTTA TTTGTTATAG TCAAAGAGAA TGACTTGTTC CTGTGCATCT ACATGAGCAT	13380
GGACCCAAA GGGTACAATT GCTCTGGAG TTGCGTGGCC GACATTAGA TTATAGACAA	13440
TCGGGATATT GCTGTCATG ATATCCAATA GTGCCTCTT ATAGTCGTCA TGGAAAGTT	13500
CATCCATAGG TTTTCCGACC AAGAGTCCAT TGATGACCGC GAATATGCCA GTGTCTTTA	13560
AAGTTAGCAA CATCTTTTG AAGTCTCTG GCTTAGGCTT TTCTTCGCTT GTTTCGAGCA	13620
AGAGGATTTT CCCTTCCCAG TCTGACAAGT CAGGGAAAAG TTTGTATTTT TGGCAGAGTT	13680
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CGAGGATTTT CCCCTCGAAC TGGGCACTTC CTTGCAACAA GTCAAAACCT GTATTTGTAT	13800
GACTGACACG AGGTGTTCCC AGGGCCGTGG GACTAAAATC AGTTCGTTCC TCATACAAA	13860
CGTCACTAGG GCGGATTCTC GAAATTCTTC CCGTCTCAAT CAATTCTTA AAGTAGTGAA	13920
GGCTATAGGC TAGCATTCT TTGTCTAATT CACAAATGTC TGCTAAAAAG GATTGACCAT	13980
AAAAAGTCTT GATTCTTAAT TTATGCAACA TGAGGTGGTT CATGGTTGTA TCCGAGAAC	14040
CAAGAAAAAT TTTTGCTTG ATAACCTTT GGAGTTGGTC ATTTTCAAAA AGATAAGGTA	14100
GCAAGCGATA GGTATCGTCT CCACCGATGG CACATAGGAT CATGTCGATG CTATCATCAG	14160
AAAAGGCATG AATCAAATCC TCTGCACGAG CTTCAAGGATG GTCCCTTGATA AAGTCTAAC	14220

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CTTTAACGA ATGGGGCAAA AAGATGGGAT TGGTCCCAGA TCCTTGAGAC GTT

14273

(2) INFORMATION FOR SEQ ID NO: 41:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 9828 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: double
(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 41:

GTGAAGTCG	GCAAAAGGTG	CAAGTGATGA	GCTCAGGTT	TTTAGCTCTT	GACATTGCC	60
TTGGCTCAGG	TGGTTATCCT	AAGGGACGTA	TCATCGAAAT	CTATGGCCCA	GAGTCATCTG	120
CTAAGACAAAC	GGTTGCCCTT	CATGCAGTTG	CACAAGCGCA	AAAAGAAGGT	GGGATTGCTG	180
CCTTTATCGA	TGCGGAACAT	GCCCTTGATC	CACCTTATGC	TGCGGCCCTT	GGTGTCAATA	240
TTGACGAATT	GCTCTTGCT	CAACCAGACT	CAGGAGAGCA	AGGTCTTGAG	ATTGCGGGAA	300
AATTGATTGA	CTCAGGTGCA	GTTGATCTTG	TCCTAGTCGA	CTCAGTTGCT	GCCCTTGTTC	360
CTCGTGCAGG	AATTGATGGA	GATATCGGAG	ATAGCCATGT	TGGTTTGCAG	GCTCGTATGA	420
TGAGCCAGGC	CATGGTAAAC	CTTGGCGCCT	CTATCAATAA	AACCAAAACA	ATTGCCATT	480
TTATCAACCA	ATTGGTGA	AAAGTGGAG	TGATGTTGG	AAATCCAGAA	ACAACACCGG	540
GCGGACGTGC	TTTGAATTTC	TATGCTTCAG	TCCGCTTGGA	TGTTCGTGGT	AATACACAAA	600
TTAAGGGAAC	TGGTGACCAA	AAAGAAACCA	ATGTCGGTAA	AGAAACTAAG	ATTAAGGTTG	660
TAAAAAAATAA	GGTAGCTCCA	CCGTTAAGG	AAGCCGTAGT	TGAAATTATG	TACGGAGAAG	720
GAATTTCTAA	GACTGGTGAG	CTTTGAAGA	TTGCAAGCGA	TTGGATATT	ATCAAAAAAG	780
CAGGGCTTG	GTATTCCTAC	AAAGATGAAA	AAATTGGCA	AGGTCTGAG	AATGCTAAGA	840
AATACTTGGC	AGAGCACCCA	GAAATCTTG	ATGAAATTGA	TAAGCAAGTC	CGTTCTAAAT	900
TTGGCTTGAT	TGATGGGAA	GAAGTTTCAG	AACAAGATAC	TGAAAACAAA	AAAGATGAGC	960
CAAAGAAAGA	AGAACGAGTG	AATGAAGAAG	TTCCGCTTGA	CTTAGGGAT	GAACTTGAAA	1020
TCGAAATTGA	AGAATAAGCT	GTAAACCGAG	TGGAGAAATC	CGCTACTTTT	TCGATTTTG	1080
ATTCAAGTTT	TTAGATTATA	TATAGTAGCT	TGAAATAAGA	TATGAACAAC	TCTATTAGGA	1140
AAGTCAAATT	AATTCTAGA	AATGTTTAG	CAGCTACAGC	GTACTATTCC	AAACTCAACC	1200
AACTATAATA	GATCGAAACT	AGAATAGTAC	ATATCTACTT	CTAAAACATT	GTTAAAATC	1260
GATTTGACTT	TCCTTATTTG	ATTCGGCTAT	ATATAGTTG	CTGTTCTTG	TCGCTCCTCT	1320
GGAAAGCTGA	TATAATAGCT	TTATGAATAA	AAAACGAACA	GTGGACCTGA	TACATGGTCC	1380

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GATTCTTCCC TCGCTCTTAA GCTTCACCTT TCCAATTTG CTATCAAATA TTTTTCAACA	1440
GCTCTATAAC ACTGCTGATG TCTTGATTGT TGGACGATTT CTTGGTCAAG AATCCTTGGC	1500
TGCAGTAGGA GCGACGACAG CGATTTTGA CCTGATTGTA GGTTTACAC TTGGTGTGG	1560
CAATGGCATG GGGATTGTCA TTGCTCGTTA TTATGGGCT CGGAATTTC A CTAAAATCAA	1620
GGAAGCAGTA GCAGCCACCT GGATTTAGG TGCTCTTTG AGCATTCTAG TTATGTTGCT	1680
GGGCTTCTT GGCTTGTATC CTCTCTTGCA ATACTTAGAT ACTCCTGCAG AAATTCTCC	1740
TCAATCTTAT CAATATATTT CTATGATTGT GACCTGTGTA GGTGTCAGCT TTGCTTATAA	1800
TCTTTTGCA GGCTTGTGC GGTCTATTGG TGACAGTCTA GCAGCCCTGG GATTCTGAT	1860
TTTCTCTGCC TTGGTTAACG TGTTCTGGA TCTCTATTT ATTACGCAAT TGCACTGGG	1920
AGTTCAATCC GCAGGACTTG CTACCATTTAT TTGCAAGGT TTATCAGCGG TTCTCTGCTT	1980
TTATTATATT CGTAAAAGTG TGCCAGAACT CTTGCCACAG TTTAAACATT TCAAATGGGA	2040
CAAAGCTTG TACCGGGATC TCTTGGAGCA AGGTTGGCT ATGGGCTTGA TGAGTTCAAT	2100
TGTATCTATC GGCACTGTGA TTTTACAGTT TTCTGTTAAT ACATTTGGTG CAGTGAATTAT	2160
TAGTCCCCAG ACGCAGCTC GACGCATTAT GACCTTTGCC CTTCTCCCTA TGACCGCTAT	2220
TTCTGCATCA ATGACGACCT TTGCTCTCA GAATCTAGGA GCTAAGCGAC CTGACCGTAT	2280
TGTTCAAGGT CTTCGAATCG GCAGTCGTTT AAGTATATCC TGGGCAGTTT TTGTTTGAT	2340
TTTCCCTTTT TTTGCCAGTC CAGCTTTGGT TTCCCTCTTG GCTAGTCGA CAGATGGTA	2400
CTTGATAGAA AATGGAAGTC TCTATCTGCA AATCAGTTCA ACCTTTTATC CCATTTGAG	2460
CCTCTGTTG ATTTATCGCA ATTGCTTGCA GGGCTTGGGG CAAAAGATCC TTCCCTCTAGT	2520
TTCTAGCTTT ATTGAACCAA TCGGAAAAAT CGTTTTGTG GTTTGATTA TTCCCTGGC	2580
AGGATATAAG GGTGTTATCC TTGTAACCC TCTTATCTGG GTTGCATGA CAGTTCAACT	2640
GTACTTCTCA TTATTCCGTC ATCCCTTGAT AAAAGAAGGC AAGGCAATCT TGGCAACCAA	2700
AGTGCATCC TAGTTGGATT TACTGAATAA AATCCATTTC CTCTAGTGAA AATCGAAAAA	2760
ACTTGTGTTTC TCTTCTTAGG TTGTTGTG AAAATAGTTT AACAGACTTT TGACTTCTT	2820
TATATGATAT AATAAAGTAT AGTATTTATG AAAAGGACAT ATAGAGACTG TAAAAATATA	2880
CTTTGAAAAA TCTTTTAGT CTGGGGTGTG ATTGTAGATA GAATGCGAC CTTGTCAGTC	2940
CTATTTACAG TGTCAAAATA GTGCCTTTG AAGTCTATC TACAAGCCTA ATCGTGACTA	3000
AGATTGTCTT CTTGTAAGG TAGAAATAAA GGAGTTCTG GTTCTGGATT GTAAAAATG	3060
AGTTGTTTA ATTGATAAGG AGTGAATAT GGAAATTAAT GTGAGTAAAT TAAGAACAGA	3120

400	
TTTGCCTCAA GTCGGCGTGC AACCATAAG GCAAGTACAC GCACACTCAA CTGGGAATCC	3180
GCATTCAACC GTACAGAACG AAGCGGATTA TCACTGGCGG AAAGACCCAG AATTAGGTTT	3240
TTTCTCGCAC ATTGTGAAA ACGGTGCAT CATGCAGGTAA GGACCTGTTG ATAATGGTGC	3300
CTGGGACGTT GGGGGCGGTT GGAATGCTGA GACCTATGCA GCGGTTGAAC TGATTGAAAG	3360
CCATTCAACC AAAGAAGAGT TCATGACGGA CTACCGCCTT TATATCGAAC TCTTACGCAA	3420
TCTAGCAGAT GAAGCAGGTT TGCCGAAAAC GCTTGATACA GGGAGTTAG CTGGAATTAA	3480
AACGCACGAG TATTGCACGA ATAACCAACC AAACAACCAC TCAGACCACG TTGACCCCTTA	3540
TCCATATCCTT GCTAAATGGG GCATAGCCG TGAGCAGTTT AAGCATGATA TTGAGAACGG	3600
CTTGACGATT GAAACAGGCT GGCAGAAGAA TGACACTGGC TACTGGTACG TACATTGAGA	3660
CGGCTCTTAT CCAAAAGACA AGTTTGAGAA AATCAATGGC ACTTGGTACT ACTTTGACAG	3720
TTCAGGCTAT ATGCTTGCAG ACCGCTGGAG GAAGCACACA GACGGCAACT GGTACTGGTT	3780
CGACAACCTCA GGCAGAAATGG CTACAGGCTG GAAGAAAATC GCTGATAAGT GGTACTATT	3840
CAACGAAGAA GGTGCCATGA AGACAGGCTG GGTCAAGTAC AAGGACACTT GGTACTACTT	3900
AGACGCTAAA GAAGGGCCA TGGTATCAAA TGCCTTTATC CAGTCAGCGG ACGGAACAGG	3960
CTGGTACTAC CTCAAACCAAG ACAGAACACT GGCAAGACAAG CCAGAATTCA CAGTAGAGCC	4020
AGATGCCCTG ATTACAGTAA AATAATAATG GAATGTCTTT CAAATCAGAA CAGGGCATAT	4080
TATTAGGTCT TGAAAAGCT TAATAGTATG CGTTTCTTG TGAGATATT TCCTTCAATT	4140
TTGCTACTAT ATTAAACAAA AATCAAAAG CAAACTAGAA AGTTATGCTC AAATAAAATC	4200
TAAATTGAC AATGAAACCC GAGTCGGATA GCTTTAAGTA CTGTTTGAG GTTGAAGATA	4260
CGATTTTGA TAGGAACCTCA TCAATTAGT ATTGTTAAGC AGCATCAATA AATTGCTTCC	4320
TTGTTTGTCT ATAATTTTTT TATTTAAAAA ATTATGACma GAGTGTGCTA TTCTTTTAT	4380
GAGAGGTGTA TGAATATGAT AAATGTATGT GATAATGTA TGTGATGTTG GAAAAAGAAT	4440
AAAAGAACTT AGAATATCTT CAAATCTTAC TCAAGATAAG ATTGCTGAGT ATTGCTCTT	4500
GAATCAAAGC ATGATTGCCA AAATGGAAAA AGGTGAAAGG AATATCACCA ATGGATTAA	4560
GTAATAAGC TTCAAACTT AGAAAAAGT TGGGAGCTGA TGTTGAATCG CCGATAGATA	4620
TTTTAAATT GGTACAAAAG ATAGAAAATT TGACGCTGGT ATTTTATGGA CTCGGAAAGA	4680
ATATTAGCGG AGTCTGTTAT AAAGGAACCTC AGTTCAGTCT CATTGAGTC AATTCAAGACA	4740
TGCCATTAGG AAGGTAAGA TTTTCTTTAG CACATGGACT GTATCATCTT TATTATGATG	4800
AGGTGAAGAA GAGTCAGTC AGTCTTATCT TGATTGGTGA AGGAGATGAA ACTGAAAGAA	4860
AAGCGGATCA GTTTGCTTCT TATTTTTAA TTTTCCCATC TTCAGTGTAT AGGATGGTTG	4920

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AGGAAATCAG AGAAAATGCC AATAGAACTC ATCTTGAAGT AGAAGATATT ATAAAATTGG	4980
GTCAGTTTA TGGTATCACT CATAAAGCTA TGTTATATAG ATTGAGGAAT GATGGATACC	5040
TTGATGCAGA AGAAATTAAA AATATGGATA TTAGTGTAT AGAGACAGCT TCAAGATTAG	5100
GCTATGATAC AAGTTTATAT CGTCCTTTGT CAGAAAGTAA AAAAGAAATG GCATTAGGAT	5160
AATATATTAA TTCAACTGAA CAACTTTAG AAAATAACAG AATTCGCAA GGGAAAGTATG	5220
AGGAACATGTT ACTAGATGCT TTCAGATATG ATATTGTATA TGGGCTAGAT GAAGAGGGGG	5280
GAGTTGTCGT TTGACTAGTC GTGTATTTAT TGATGCAGAT TGTATTCAG TATTTTATG	5340
GGTTGGCACT GAACATCTTT TAGAAAAGCT CTATTTGGGT AAAATTGTAA TTCCACAAGA	5400
GGTGTATGAT GAAATCAATA TACCTACAAT TCCCCATTAA AAATCTAGGA TAGATCAGTT	5460
GGTAGCTAAG GGTTCACTGAG CATAGACATT GGAACGTGAG AATACGCATT	5520
ATATAGAGAT TTAACAAGAA ATCATGATAG TAACAAGATT ATTGGTAAGG GAGAAGGGC	5580
ATCTATTTCC TTAGCGAAAA AGCATAATGG GATATTAGGA AGTAATAACC TAAGAGATGT	5640
TAAATCATAT GTAGAAGAAT TTCTTTAGA ATATATGACA ACAGGAGATA TACTGATTGA	5700
AGCGTTTAAA GCGTAATTAA TTACTGAATA AGAGGGCAAT CATATCTGGA ATAATATGCT	5760
TAAAAAGAGA AGGAAAATTG GTGCAAATTC ATTTTCAGAC TATCTTCGTG GAAGTATTCA	5820
TCAAAATAGA CAAAAATAAA TTTGGATAAA TCGAACTCAC TATTCAGGAG GCATATGAGC	5880
AATTGGAAAA AGAAAAGTGT CAAATTGAGC CTATAGGAGT AGAAGTGAAA TAGTAAGTCC	5940
TGCATAGTGG ATGAGAGAAA AGTCTCCCTT GAAGTTTCC TGAACATATCA GTCGCATGTC	6000
AAACGATATG TAGGTAATG TGAGAGGGGA TAGCGAGTAG TTTTTGGTTA TTTTATCAA	6060
AAACTTATAT TTATTATATAC CGAATGATAA AATATAATAA AAATGATAGA ATAAGGAAAA	6120
AACATGAATG TCAAAAAGAT AATGTCAATT TTTCATCCTT TTATGTTGA TGTCAGTATT	6180
GAGGAACGTGA CTTTGACTTT ACCAACATCAGT TTGTAAAAA GGTTTGAGTA TACTCAAATG	6240
ACTTTTCATA AGGAATCATT TTATTGATT AAAGAAAAGA GAAGGGGGAG TTTGAGTTCA	6300
TTTGTACTC AGGCTCGCAC TATGGGTGAA AAAGCCAATA TGGATGTTGT TTTGGTGT	6360
TCGAAGTTAT CACACAGTGA AAAAAAGCAA TTACTTCAG CTAAGATTCC GTTTGTAGAC	6420
TTTAAGGGAA ACCTCTTCTT CCCTCCATTG GGACTAGTAC TCAATGCGAA TGATACTGAA	6480
GTCCCTAAGG AATTAACACC TAGCGAACAA TTAACGTGGA TTGCTTTTT ATTGACAAAA	6540
GGTCAAAAG TAGTAGATGT TGATTTGCTT TCACAAGTCA CTGGACTTCC AAACCAACAA	6600
ATTTATAGGT GTTGAGGAC TTTAAAGCT TTATATTGGT TAAACAAAGCA AAATAAGCTT	6660

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TACACATATA CGGTGTCAAA GAAAGAATTA TTCTTAAAAT CCGTGTATG TTTATTTAAT	6720
CCCATCAAAA AACGGATTT ATTGCCAGAT GGCGATATAA AGCAGATAAA ATCTGTTCT	6780
AACCTTCTAT ATGGTGTGC TTATGCTTG TCGCATTCAA CTTTTTTAGC TGAAACGGAT	6840
GAAAATATTA GCTATGTCAT ATGGCAGAGA AAATTCAATC AGTTATCCTT GCCACTTCT	6900
CAGCATGTT TAAAATGAAA GATGCTAGAG ATATGGAAAT ATCGTCCTT TGTATCTGAG	6960
TTTTGGAATG ATTTTAAAAA TAATCATGAT AAACAATTG TAGATCCGAT TTCTCTTAT	7020
TTGACCTTAA AAGATGATGA TGACCCACGT ATAGAGGAAG AGAGTGAAGC ACTAGAAAAT	7080
ATGATATTAC AGTATCTGGG AGAAGATGAT GCCAGCTAAT ACGAAAGTTA TTTTTCAAGA	7140
AATGTTGCG GATTTTCAGA ACTATTATGT TCTGATTGGG GGAAC TGCTA CCTCTATCGT	7200
ATTGGATTGCG CAAGGATTAA AAAGTCGCAC AACAAAAGAT TATGATATGG TCATCATTGA	7260
TGAAGTAAAA AATAAGGAAT TTTATTA CTTGAATCAT TTTTTAGAAT TGGGAGAGTA	7320
TCAAGGAAGT CAGAAAGATG AGAAAGCGCA GCTTTTCGA TTTACAACAA CTAATCCTGA	7380
GTTTCCCTCT ATGATTGAAC TATTTAGTAT CTTACCGAA TATCCATTA AGAAGGACGG	7440
TCGAGAAATT CCCTACATT TTGACCAAGA TGCTAGTTA TCAGCCTTAT TATTGGATGA	7500
AGATTATTAT AATATATTGG TGCAATGAAAA AGAAACCATT CAGGGTATT CGGTATTGAG	7560
TAATTGTTGGT TTATTA CTTCT CGAAAATCTC TTCAAAACCAC GTCAAGCTTCC ATCTACAACC	7620
TCAAAACAGT GTTTGAGCA GCCTGCAGCT AGCTTCCAG TTTGCTCTTT GATTTTCATT	7680
GAGTATTAAT TATTTTTAAG GCTAAAGCTT GGCTGGATAT GAGGGAGCGC TCTGCCACAG	7740
GTGCTCAAGG TTTAAGTAAG TCCATTAAAA AGCATTGAA TGACCTTACC CGTTGACAG	7800
CTTCCTTGCT AGGAGATGAA AAGTTATCGG CTATAACATC AAGTAGTGCG GTAAAAGCAG	7860
ACATGCACCG CTTTGTGATA GAATTAGAGC CTGTGAAGTC AACTATTCTT CAAAATAATG	7920
ACATTTCATT GGATCAAAAT GAAATTTTG AAATTCTGAA AAATTTCTC GATGGTTAAA	7980
ATAATTGTAG CGAGATGGCT ATATTGAATT CGTCTATATC TGAAACTAG AAAAACCTTC	8040
AATTTCAAGA GAAAATGAAG TCAATCTTCC CACAATCAA CGTATAGTAT CAAGGTTTT	8100
CAAGACCTGA TATTATGCGT TTTTGCTTT TCAAAACTTT TTGCCAGTC TTGCTTTTA	8160
TCCTCTAGTC ACTTGATTTG TTTCAAGGTGG TTTTTAGTA TAGTAGAATG AACGAGAAC	8220
AGGACAAATT GATCAGGACA GTCAAATCGA TTTCTAACAA TGTTTAGAA GCAGAAAGTGT	8280
ACTATTCTAG TTTCAATCTA CTATAGTTAA ATCTGCGGTC AAGTCTACTG GTGAATCTAT	8340
GATTGTAATA CTCTCCAAA ATCTCATCAA CCACGTCAGT CTTGCCCTGAG AGTCTGTATC	8400
TTACTGACCA AGCTAGTGAT GGATTTAGAA TAGGTGATTT GGAGCGTCCT ATTAGCTAGG	8460

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AAATGCTGCT CATA GTC CTT TGCTGAGGCT AGGGTGTTC AACATTCAAC ACTCAACTGG	8520
TTGATCTAGT TGATAGGAAG GGAGTTACTA TAAAATACTC AGGCTTCCAT CATATTTTT	8580
GAAACGATTG TGTAAATCAAAT ATGTACCAAT ATTGTAGTAT TGGTACAGAA GATGTTGTGA	8640
ATGGATAAAAT ATATCATAAC TGCTATCTCA AAAAGATTTC ATATGTCTGT GCATATATAA	8700
TAGACTTCCCT GCAAACACTAG AATCCTAGTT CATGATTGAT AATACCAGCA ATCAAATTCA	8760
TTCGTAATCC AAAGCGTTA CGATGATTTC GATAGTTGT TGAAAACATT TTAAACGTTT	8820
CTACTTTGGC AAAGATGTTC TCAACCTTGC TTCTCTCCTT AGATAGCGCA TGGTTATAGG	8880
CTTTATCTTC AGCTGTTAGC GGCTTGAGTT TGCTGGATT ACGTGGAGTT TGTGCTTGAG	8940
GACATATCTT CATGAGCCCT TGATAACCAC TGTCAGCCAA GATTTAACCA GCTTGTCCGA	9000
TATTTCTGCA ACTCATTTTG AACAACCTCA TATCATGACA ATAGTTACACA GTGATATCCA	9060
AAGAAACAAT TCTCCCTTGA CTTGTGACAA TCGCTTGAGC CTTCATAGCG TGAAATTCT	9120
TTTTACCAGA ATCATTGCT AATTCTTTT TTAGGGCGAT TGATTTTAC TTCCGTCGCA	9180
TCAATCATTA CCGTGTCTC AGAACTAAGA GGAGTTCTTG AAATCGAAC ACCACTTTGA	9240
ACAAGAGTTA CTTCAACCCA TTGGCTCCGA CGGATTAAGT TGCTTTCTG AATACCAAAA	9300
TCAGCCCAA TTTCTTCATA AGTGCCTAT TCTAGGCTTA ATTTAGGTTT TCGTCCACCT	9360
TTTGCCTGTT TAAGTTGATA AGCTGTTTT AATACAGCTA ACATCTCTT AAAAGTCGTG	9420
CGCTGAACAC CAACAAGACG CTTAAATCGT GTATCAGTTA ATTTGTTACT TGCTTCATAA	9480
TTTCGCAAGGG AGTCTATTGA CTCTTTGGTA GGTGTCAATG TTTTTTCAT CTATCCCGAG	9540
AATTATTTTC CCGCCATTG TATTTGCAAA TCCTGAGTAG GTTTCCCAGA AAGACTCTGG	9600
AAGATTGTTT TTAGCTTTT TGTATTCTAA ATCAACCCCT TCAAATTTTA AGTCCCATATT	9660
TTTCCTTAC ATCTGTTTT TGTGGTTCTG GTATTTCTC AAGTTGAGTG ATAATATAGC	9720
GAATTGAATT TCGAGAGTTT TTACTCAGTT AATTTCTTTT TTAACCCACT TTAATTGCTT	9780
TTTTAACACG GGTTAAAAAA GAAATTAAAG TGGGTTAATT TTTCTTGA	9828

(2) INFORMATION FOR SEQ ID NO: 42:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 3369 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 42:

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CCCGCAAAGA TATTTTGAA CAAGAGTTG GACGTGAGGT CCGTGGCTAT AATAAAGTAG	60
AAGTTGACGA GTTTTAGAC GATGTCATCA AGGACTATGA AACCTATGCT GCCTTGGTCA	120
AGTCACCTCG TCAGGAAATT GCCGATTGAA AGGAAGAAATT AACTCGTAAA CCGAAACCTT	180
CACCAAGTTCA AGCAGAACCC CTTGAAGCGG CAATTACAAG TTCTATGACC AATTTTGATA	240
TTTTGAAACG CCTGAATAGA TTGGAAAAAG AAGTTTTGG TAAACAAATT TTAGATAACT	300
CAGATTTTA AGTAGTTATT TGAGATGTGC AATTTTGGA TAATCGCGT AGGAGAATTG	360
TTTCTCATGA GGAAAGTCCA TGCTAGCACA GGCTGTGATG CCTGTAGTGT TTGTGCTAGG	420
CGAAACCATA AGCCTAGGGA CGAGAAATCG TTACGGCAGT TGAAATGGCT AAGTCCTGG	480
ATAGGCCAGA GTAGGCTTGA AAGTGCCACA GTGACGGAGT CTTTCTGGAA ACAGAGAGAG	540
TGAAACGCGG TAAACCCCTC AAGCTAGCAA CCCAAATTGTT GGTCGGGCA TGGAGTACGC	600
GGAAACGAAC GTAGTATTCT GACTGCTATC AGCTAGAGCT GTTACTGGTA GACAGATGAT	660
TATCGAAGGA AGTGGCTCTA GTCACTTCTG GAACAAAACA TGGCTTATAG AAAATTGCAT	720
ATAGGTTGGG GCTCAGAAAT TTTCTCAACC TCATTTTTA AAGTGGACAT ATAGAAAGGT	780
CTTGCAGAC TGTAACATGA AAAAGAATT TAATTTAATT GCAACTGTGG CAGCAGGGCT	840
TGAGGCTGTC GTTGGCTGTG AAGTGGAGA GTTGGGCTAC GATTGTCAAGG TTGAAAATGG	900
ACGTGTTCGT TTTCAAGGAG ACGTGAGAGC TATTATCGAA ACCAACCTTT GGCTTGGGC	960
AGCAGATGCT ATCAAAMTA TCCTAGGAAAC GTTCCCACCT AAGACTTTG AAGACCTATT	1020
TCAGGGAGTT TTCCCTTGG ATTGGAAAAA TTATTTACCA CTTGGAGCTC GGTTCCCGAT	1080
TTCAAAAGCT AAATGTGTTA AGTCCAAACT TCACAATGAG CCCAGTGTTC AGGCTATTTC	1140
TAAGAAAGCT GTTGTCAAGA AATTGCGAAAC ACACATATGCT CGCCCCAGAAG GGGTTCTCT	1200
GATGGAGAAT GGCCCAGAGT TTAAGATTGA GGTCTCTATT CTCAAAGATG TGGCAACTGT	1260
CATGATTGAT ACGACCGGGT CTAGCCTCTT TAAACGTGGT TATCGTACCC AAAAAGGTGG	1320
CGCTCCTATC AAGGAAAATA TGCGAGCAGC CATTTCACAA CTTTCTAACT GGTATCCAGA	1380
CAAGCCTTG ATTGATCCGA CCTGTGGTTC GGGGACTTTTC TGTTATGAGG CAGTTATGAT	1440
TGCTAGAAAG ATGGCGCCAG GTCTTCGTG CTCTTTGCA TTTGAGGAAT GGAACGGAT	1500
CAGCGATCGC TTGATTCAAG AAGTGGCAC AGAAGCGGCT AAAAAGTAG ACCGTGAGCT	1560
TGAGCTGGAT ATCATGGGCT GTGATATTGA TGCTCCATG GTGAAATTG CTAAGCCAA	1620
TGCTCAGGTA GCTGGTGTG CAGGAGACAT TACTTTAAG CAGATGGCGG TGCAGGATTT	1680
ACGTTCCGAT AAAATCAATG GAGTAATCAT TTCCAATCCG CCTTATGGTG AACGTTGTGTC	1740
AGATGATGCA GGGGTGACCA AGCTCTATGC TGAGATGGGG CAAGTATTG CACCGCTGAA	1800

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AACTTGGAGC AAATTATCC TGACTAGTGA TGAAGCTTT GAAAGCAAGT ATGGTAGCCA	1860
AGCAGATAAG AAGCGTAAGT TATACAACGG AACCTTGAAA GTGGATCTAT ATCAATATT	1920
TGGTCAGCGT GTCAAACGGC AAGAGGTAAA ATAGAAAGGG ATACTCATGA GTAAAAAAAG	1980
ACGAAATCGT CATAAAAAAG AAGGTCAAGA ACCGCAATT GATTTGATG AAGCAAAAGA	2040
GCTAACAGTT GGTCAGCTA TTCTGAAAAA TGAAGAAGTG GAATCAGGAG TCTTGCTGA	2100
GGATTCCATT TTGGACAAAGT ATGTTAAGCA ACACAGAGAT GAAATTGAGG CGGATAAGTT	2160
TGCGACTCGT CAATACAAAA AAGAGGAGTT CGTTGAAACT CAGAGTCTGG ATGATTTAAT	2220
TCAAGAGATG CGTGAGGCTG TAGAGAAGTC AGAACCTTCT TCAGGAGGAAG TTCCATCTTC	2280
TGAAGACATC TTACTACCCCT TGCCCTCTGGA CGATGAGGAG CAAGGCTTGG ATCCCTCTATT	2340
GCTAGATGAT GAAAATCCAA CAGAAATGAC TGAAGAAGTG GAAGAGGAGC AAAACCTTTC	2400
TCGTCTGGAT CAAGAGGACT CAGAAAAGAA AAGTAAAAAA GGCTTTATT TGACCGTTTT	2460
GGCGCTTGTA TCAGTAATT TAATGTCAG TGCTTATTAT GTCTACCGTC AAGTGGCTCG	2520
TTCGACTAAG GAAAATGAAA CTTCTCAATC AACTACAGCC AATCAATCGG ATGTGGATGA	2580
TTTTAATACA CTTTATGACG CCTTTTACAC AGATAGCAAT AAAACGGCTT TGAAAAATAG	2640
CCAGTTTGAT AACTGAGTC AACTCAAGAC TTTACTTGAT AAGCTGGAAG GTAGTCGTGA	2700
ACATACGCTT GCCAAATCTA AATATGATAG TCTAGCAACG CAAATCAAGG CTATTCAAGA	2760
TGTCAATGCT CAATTTGAGA ACCAGCTAT TGTGGATGGT GTGTTGGATA CCAATGCCAA	2820
AGCCAAATCG GATGCTAAAT TTACGGATAT TAAACCTGGA AATACGGAGC TTGATAAAAGT	2880
GCTAGATAAG GCTATCAGTC TTGGTAAGAG CCAGCAAACA AGTACTTCTA GCTCAAGTTC	2940
AAGTCAAACT AGCAGCTAA GTTCAAGTCA AGCAAGTTCA AATACGACTA GTGAGCCAAA	3000
ACCAAGTAGT TCAAATGAGA CTAGAAAGTAG TCGCAGTGAA GTCAATATGG GTCTCTCGAG	3060
TGCAGGGGTT GCTGTTCAAA GAACTGCCAG TCGTGTGCCC TATAATCAGT CTGCTATTGA	3120
TGATACTAAT AACTCTGCCT GGGATTTGC GGATGGTGTC TTGGAACAAA TTCTAGCGAC	3180
TTCACGTTCA CGTGGCTATA TCACTGGAGA CCAATATATC CTTGAACGTG TCAATATCGT	3240
TAACGGCAAT GTTATTAACA ACCTCTACAA GCCAGATGGA ACCTATCTCT TTACCCCTAA	3300
CTGTAAGACA GGCTACTTTG TCGGAAATGG CGCTGGTCAT GCGGATGACT TAGATTACTA	3360
AGCAGTCGG	3369

(2) INFORMATION FOR SEQ ID NO: 43:

- (i) SEQUENCE CHARACTERISTICS:
- (A) LENGTH: 9713 base pairs

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- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 43:

AAGTTTACAA TTTAAATGAA TTAACAATTT TCCCAACTAA AAGCACTCCA GTTACCGCAA	60
CGTTTGTACT GAATGTACTA AATCGCATTC CATCAACTTC ATCTGTTTCG TCAACTTGAA	120
CAGATACTAA TTGAAGATTG AATACTTCTG CTGCCATAGC TAGCTCCTCC TATTTAAATT	180
TTTGGGATTA AGTACTTTAT CCACCCCTCAT ATACTCTCTC CACCAAGTAAA ATGCAAGCAA	240
TGATACAAAA TAGATTTAAC TATTTTATAT AGCGAAAAGT TACAAATTAA TAAGAAATAA	300
TTTTTGCAATT CTTAAAGATA AAATAGGAAC TTCTAGTAAT AAATATTAAA ATAAATAAAA	360
TAATAGATAAC TATAAAATTG GGAAGTATTA ACCCCAAAAG ATTCAATATCA TCTATTAAA	420
TATCCTCTAA AGAGTAGTAT ATAAAGCCA TAATTTAAT GTTAAGTAAA ATGCAATTAA	480
ATGAAGTAAC AAATGTCAA AATATAGCCT CACCAACTTT AATCTTAACC ATCTGGTAAT	540
TAGAAGTTCC TAAAATTCA AATTGCTGAA TCTCAATCCT TTCTTGATGC GATGACAAAA	600
ATGCAATTGA AATAATATTG GCAAGTACTA TCAAAATTGG TGCTCCTACA TAGACAATAA	660
ATGCTACTTT TAGCTCTAAA TCACGTCTCAT CTTGAAATTG AGATAGTATA TTCTGAGAAA	720
TCATTTGAAA ACTACAAATT ACTAATATAG CTCCTGTAAT TGCAAGCACTG ATAGATTTA	780
TATAAGACTT ACAATATAGT AAATTCCACT TCGAAACAAT GAACATAAAA TTATTTCTAA	840
ATATAATTAT AGAAAGTAGT TTGATAAAAC ATGACTGTAT AAAAGGAGAT AATTGATAAA	900
TAATCACAAT ATCTAAGATT ACAATATTGA ATATTATCTG GGCTTCGCT AAAATTGTGC	960
TATCTGGAA AATTGTTGC AAAGAAAGCA ACCAGATAAC ACTAAACCA GCCAATAGCA	1020
GTATTCTTT TACTATTGAA AGAACATGCC TTATTTAGA ACTCTTCCTA TTTCTAATCT	1080
TCTTGACGT ATAAAAGCAA CCACCTGAA AGGCTAAAAA TGAAATCAAC ACTACTGTAA	1140
TGATACATCC AACAGCACTC GTTTGAAATT GGATATCAGG TAATATATTG TCCCCGAAAA	1200
AGTATTGTAA AAAATAATAA TAATTTGACG TAACAAATAT AGAGCATAGA TATGCAATAA	1260
AACTAATAAT CGAGGAAATG ATAAAAATCT GTCCCCCAC AAGAAATGAT AGTTGAAGGC	1320
GACTTGTCC CAACACCTCC AGAAGTTCGT AATCATCTCT AAAAATTCA ACCAACATAT	1380
TTATTATGTT AGAGAGCACA AAGAATAATG TTACTCCTCC GAATACTATC GGAAACATAA	1440
AAATTGGTTT AGGATCTGGA AGTCCGACAA ATACTTGCAG ATTATTCCTCA ACATTAATTA	1500
CCCCATTAAC AGCCAATCCC ATAACAAAC TCGAAACAAA AATTACTGGT GAAACGCCTA	1560

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ACCATTGTTT CTTATTATGT AAAAATTGAT AGTAAACTAA TCTGAGCATE TCTATTCCCTC	1620
CGTAGTTGAT TGTACCTCTA AGATTTATA CAACTCTTCC CCGCTAGGTC TATGAAGTTC	1680
TTTGAAAATT TTTCCATCTT TCAATATTAA TGACAGATCA GTTTTCGAGG CCAATTCTAT	1740
ATCGTGCCTT ACCATAATTA CACACTTAC CGCCCCACT AACTCTCTCA ATAATTCAA	1800
AATTACTTCA CGAGAACGCG TGTCTAAAGC CCCAGTTGGC TCATCAGCAA ATATTATATC	1860
ACTATCAGCA ATAACCGCTC TAGCTATAGC AACCTCTGTG TTGTTCTCCAC CAGACAGAGT	1920
TCCAACAAAA TCGTTTAAGC CAGCATTAAA CTTCATTCTT TTGAGTAAGT TTTCTACATT	1980
TTTAATAGTT AATTTTTTTT GTGATAATCG CAAAGGAAGT GCTATATTCTT CTATTACCGG	2040
CAGGGAAGGT ATTAATTGAT ATGCTTGAAA TATAAAAGAT ACTTCGTTAC GTCTTATACT	2100
TGACAATTTC GCATTCTGA TTTTATAGGG GTTGATTCCA TTTAAATTAA CTTCCCCACT	2160
TGTTGGTTCA AGCAAACAG AAATACATTT TAATAAAGTT GACTTTCCAG AACCACTAAT	2220
TCCTAGAATA CTTATAAATT CTCCCTCTCGA AGCAGAAAGA GAAACATTTC TCAGCACTTG	2280
CAACGTTTA TTATTCCTA GTAAAAATTG ATGATACAGC CCTTTCACTT TTAATATATA	2340
ATCTTATCC ATATTCTTG CTCACATCAC TTAATTTGA AAAGTGTTC ATTTCCTAAC	2400
TTATATATAT CAGTGTATCT CTTGTCATTT AAGTCATAAT GATGTGAAAC TTCAATAAAT	2460
GAAATACTA AATTGAACAG AATATCATGT ATGGAATTG AATTATCATT ATCTAAATTAA	2520
GCTGATATTT CGTCAAATAA GTACACTTTA TTATTCCTAA TCAGAGCTCT AGCTAAAGCT	2580
ATTTTTGTT TTTGACCTCC AGACAAATTAA CTACCATTTT CACCACATTG ATAATTAGT	2640
ATATCTATCT TTTCTAATTC TTCATATAGA TTTACCTTTT TTAACACCTC AATTATCTGA	2700
TCATCTGAAA AATATTCATT TTGAAATAAA GTTACGTTCT CACGAATAGT AGTGTCAAAA	2760
ATATATGGTG TCTGATCAAC TGTGGTATT GAATCTGAAC TCTTTTCCC ATGTGATAAC	2820
AAATTACAT AACCTTTTG TGCTTTAAA GAACCATTAA TTAAATTAA AATCGTTGTT	2880
TTCCCACTAC CAGAAGTTCC TGTTAATAAT ACCCTAAATG GTGACTTAAA TGAGAAGTCA	2940
ATACTTAATT TATTTCTGG TGTATAGAA TATACAACAT CTTTCATGTG TATCTCATCT	3000
ATTGATGAAG TATACAGTCC GTTATTATCA TGTTCACTGT CTATAAAATT CTTCTCTCCA	3060
CTTAAGTATT TTAAAAACGG TTTCCTTAAA TCTTTGGTTG TATTTATCTT ATTTAATGAA	3120
TAGGCAATTG ATTGTATCGG CCCTAAAATC TTATCGTTG CTAAGAAAAT ACCTATCAGT	3180
TCACTAAAAG AAAGGCTTTT ATGATAAAATT ACAAAATAAC ATCCTACAAC CAAGGAACT	3240
AGAAAGCAAA AACCTGAAAT TAGTACTGCA ACCAATTTCG AAAGAACCTC TGATCGTTTC	3300

408	
AAATTAAAAG TAGAACCTTC TAGTTTATCC AACTTTTAT CCGACAAACT AATTATTTCT	3360
TTAGTAACAG AATAAGATTT TAATGTCTTA AAACCATTAA AAATTTCTTT TATTATGTGA	3420
GTATACTCTG CATTGCTGTT AGAGTACTCA TTAGCTGAAT TAGACAACAT CTTCTTCATA	3480
AAGACAGGTA CTATAATCGG CAATGCTGAT AATAACAATAA ATATTATTGA nACTAGGAAG	3540
TTTAAATAAA GCATAAAAACT TAGAGAGACG ATGAACAACA ATATTGAAGA AATTATTTCA	3600
AAAATTGTC TAAAATAGTT TTCTTCGATT AATCTCAAAT CATTGACAA AACTGAAATA	3660
ATAGATGAGT AATCTTAAAC CATTTCAGAA GAAAGATACT GTTCTCTAAA ATATCCTTGT	3720
TTAATTTTA CATTATATC TTTAGTTATT GATGCTTCCG TTACTTCTAA ATAGTAATT	3780
GATATATAGA TTGCTGACCA ACCCAGAATA CTTATAGCAC CAAATCTTAG AACGTCAGAA	3840
AATGAGGAAG TCTGATTTAA ACTACCTGCA TATACAATAA TTCCTGAGAG CAAGACACCA	3900
TTAAACGAAG ATAGAAAAT TAAAATCCCC ATTAATATAA GTTTAGTCTT TTTTATAAAT	3960
TTTAAATAAT TCATAAGTTA TTCTTCCCCTA CTTCTTCAAA GAAATAATTT AAAGTATCAA	4020
TCATTAAGAG AACATCTGAT GGAGTAAAAC CTCCATGACC AGCTGCTTG TTTAAATACA	4080
ACAAACTTTT AACTCCAATA GAATTTAATT TCTTGACCA CTCTATCACT TCGTTATTAT	4140
TAATATATGG GTCTTCTCA CCCAAATAT TAATCTAAAC AGTATTTGAG TCTCGTGCCT	4200
TTTCAATATT TTGCTATAGGC GAATATGACT TTATATAAGC CTTTACTTCA GGGTCTCTAA	4260
TATCTCCCCTA CTCTGCTATT TCGGTCTTAG AAAGAGGATC ATTTGGATTC TGAAGTGTAT	4320
CATAAGGATT TATAATGGC GAAAATAAGA GAATGCTTG CAATAAATTT TTTTCCTCGT	4380
TCAACACCGC ACCAGCAATT ATTCCACCTG CACTAGAACT TATTAACCT AATCGCTTAC	4440
TGTCAATTAC ATCATTTTCC CTTAAATAAT TTACTCCCTC AATAAAATCT CTGATAGAAT	4500
TCCATTTGTT TAACGCCCTT CCTGAGCGAT ACCATTCACC ACCAAATAG CCTCCACCTC	4560
TTACATGAAC TATAGCATAA ATAAAACCTG CATCTATTAT AGATAACATA ATTTCATCTA	4620
AATCAGAATT ATCATTCTTA CCATAAGCCC CATAGACACT TAGAATACAT TTTTTCTCTC	4680
TTGGGAGCTC ATCCGTATCT TCACTTTCC AAAATAAGA AATCGGTATG CTTACATCAT	4740
AACTGTCTTT TTTAGTCCAA ATCACCTTAG AAAATATTT AGTATTATTC GATTTTATGA	4800
TGGGTCTTCA AAATTCAGTT TTTAATGTAT TTTCTATTAA ATCAAAACTA AGTATTTTT	4860
CGTAAAAAGT TCTCCTCTCT AAAACAGAA GAACACGATC AGAAAATGAA TTTTCATAAA	4920
GTGTTGTCTT TTCATCAAAT GTTATCTTAT TAACACTCAA CTCCCTCAA CTATTATTT	4980
TAAATGTAGC AAGATAAAAG ACGGAATTGCG CTGCGTTGA ACAGTCTAA AGGATATAAC	5040
GTCCTATACA GTGAACTCTT CTAGCCCTAT CTTGATATGG TATAGTAATA GAAACTCTGT	5100

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CTCCCGAAGA AGTTTCCCTT AGAATTAGTT GATCTTCAGTT	GAAGAGAGCC	5160
CAAGAAAGTA CTGTGCTTT TCTGTACTAA ATAGAGCGAT	ATCTCTAGGT	5220
CCGTTTCTGT GTAAAGAGTGT CTAACAAAAC CCGTCCGGTC	GAAACTGTAT	5280
TGCCTTCTGT AAAGTCTACT GACTTTACAA ACAATTATT	GCTATCAATG	5340
TTAACGAAA AGAGCATTG TTTTCTCAA ACAGTTCTC	TTCTGTAAAG	5400
ATTTATAGAA TAACCTTACTT GGCTCCCCT ACTCTTTGGA	GCGAGTATAC	5460
ATTTACCCAA ATAGAACGAA CTTTCTACTG AATATCTTC	AATGATAAAT	5520
TAGTATATT TTCTTATTCCA ATTAAATTAG TCGTACGCAG	TGAGGATACA	5580
TATAACTCTC ATCAGATGAA ATCCTAACAT CCTGTAAGAT	ACTATCATCT	5640
ATTTTCCAC ATCAAAGACA ATTTTAAGTG AATTGAAATT	GTCTAAACTG	5700
CCTTAGGAAT CCAGTCATTA TCTTCGACAT ACCATTCTT	TATTACACCA	5760
TACTCCAATT ATCAAATTGG TACCAATATC GCCCTCTCCT	AAATATCAA	5820
TTTTTAATTCT CTGAAATGAT GAAGAGATAG ACCTCTTATA	GTGTGTTTT	5880
TATTTAAAAA TATTCATTA CTCTGATTCA CAAGTATGAC	CCCTTAATAA	5940
ATATTATATT TGAGGAAGAA TCGTCAATT ATTATCCATT	ATTGATACCA	6000
AACACCCGCA AATCCCGAAG CAATATCTGT TGTTATCTT	AAACCATTAT	6060
AAACAAATCCT TCTTCATTA CACACAAATA TCTATAAAGT	TGTTCAATTA	6120
TCCTGAAAAG TTATCATCGA TATCACTATA TATATTATTA	GCAACTTCAA	6180
TCCGTAAAT AACCTGGTA ATACACAAAA AACTACATCA	GACCACAAAA	6240
TAAATATTTT AAGTATTGTC TTGACAAGAT TTCTTTATTT	CTATTAATAA	6300
GCCAGCACTT CCAGTTGCTA GATATGGTAG TAATCTATGA	CCTTGGCTGT	6360
ATTATTACTA TCTACTTTAT AAGCAACTAA TTCTTTATCT	ACAGCCAATT	6420
TTTATAGATA CTTCACCAAG TTAATTATA AGCTTCACCG	AAGAGCCAAG	6480
GTGACCATAT AGTAATCCAC CAAAATTCTC ATAAGGATCG	TTACTCTGAA	6540
GCCAACTTTA CAAAAGTTT CTGGATTTTC TATATAATT	AAAGTATATT	6600
AATTAGTATT TCTTCTCTA GTTTATTATC AATTCCCCCT	TTACTAAGAA	6660
AACCAGTAAA ATTCCAGCCT GCCCACTATA TAAATTTTA	TTTGTGAAT	6720
CTCTATAAAA TGAGTTGTAAC AAAGTTAAC TGCCCGATCT	ATCTCCCCAA	6780
GAGCCAGATT GTACCAATT TACCATCAA AAGACCAGAA	AGGGACCGATT	6840

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ATTTACTGCC TCATTAATAA CCTGTGTTCG AATCTCATAA TAGTCATCAA ACTTGAAATT	6900
TTTTACTTTC TTAGCTAGTT GTTGATAACT CCAAAGGATA GCTAAATCTG AAAACGCAAT	6960
TCCCTGATTA AAATTCAAGAC CATAATAATG AACTGGGAAG AATCTTGATT GAAATTCTTT	7020
ACGCCACTGT CCATAAGTTA CGCTAAACCC TCTCAATAAT TTTATAATAA AATCTTGAT	7080
ATCTTGCTCA CTCTCGATAG TTCTAATCTC ATGCATGGGT TTTAAAACCTT TTTTCCTGGA	7140
AATATTCTCA ATCTGTGGAC ATTPTAGAATC TAGATATGAC AATAAACTTT CTACATAATC	7200
TATATGTTCT CTTGTATAAC CCAAAGACTC AAATAGTTTT TTTCCCTCTA TCCTGGTTG	7260
ACTTACATAG TTGTATGTCA AATCCGATGT AGTTACTAGT GGCGATGTATA AATAATGAGC	7320
TATTTGTCTA ATACCATAACC AATCTATCTC ACTGGGAAGT GTTTCTCGCC ATGCTCTAAA	7380
ACCAGGGGCT GCAACTTTAT GTACAACCTT TTCATCATTT GAAAAGACAG CCTGTTCCC	7440
GTCTATTATA CTAATCTCAT CTTCATCCTT AACCAAGATA TTTCTAAAT GTAAATCTTG	7500
ATGATATACA TTTTCAGAAT GAAACTTATT CGTTAAATCG ATGAGTTTT CTACTATCTT	7560
TGAAACTCTC AATAGATAAT CTTGGTCTT ATCAACAACT TCATATAAAG GAAAATTATT	7620
GGTAAACCCAT CTATTTAGTG GAACGCCCTT CATAATGTTCA ATTCTTAAGA AGGTGTGCTC	7680
CCAGATCTTA CCGTGCCAGT ATATTTAGG CGTCTCACTC CATTCAATTAA GAATTTTAG	7740
TGCTTTGCAC TCCGAAGCTA ATTCTCTGA AGAATAAGTA CCATCAAATC CTAGACCTGT	7800
ATACGGTCTA GCCTCTTTA AAATTATTTT TTTCCCATCT TCTTTTAGCC TAGCATTATA	7860
TATCCCACCA CTGTTGAAA ATCTAATTGC ATTATCTATA ATAAAGGGAA AGTCTCCCTG	7920
TTTTTTATCT TTCTGTCAA GCCATTTATT CAAAAAGTC GGGGGCACTA TACCTTTGG	7980
AATTTAAAT ACTGGTAAAC GTTCATCTTT ACAACCTCA TCGCCAACAA TTAATTCTAC	8040
AATAGCAACC TTCTTTCTAT CATCCCTTGA CGGCCTAAAC ACACCATAACC TCAGATATAT	8100
TGGTGCTTCA TCCCAACGTT TATCGTTAA AATATATGGC CCATTATATT GCTTTAAGGC	8160
ACTTTCTAAC CTTTGCAAAA CCGACTCTAA TTCATTTGA TTTGGATAAC ATGTAATAAA	8220
TTTACCCAGAA AATCCTCGAC TAACCAATT CCCGTTTCGC ATGATAAAATT TGTCTTCTGT	8280
ACTAAGATGT TTAAATGGAA TTGCGATTTC ATGGCAAATT TTTGCTACAT CTTGTAACAA	8340
TTCATGTGAA CTGTTTAACT CTGAACATAAT GTGTATTTC CACCCCTGTC TTTCAACAAA	8400
TTTTCCAATA GGGTATTGAT AAACCCACTC ATCATTATTC ATTACTTCGT GCCAATTAAA	8460
AGGCAGACTT ACTTGGTACT TTATGCTAGT ATCTGTACTA TAATCATTAT TAGTGAAAAA	8520
GAAAGGATGC TCCAAATTGA AATTATAATC CATAACAAAA TCTCCAAGAA ATTTTATCAA	8580
ACTTAATATA TCTATAGCTA GACAGACTTA TTTAAATAAA AAGGGAGAAT CCTTTGGATT	8640

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CTCCCCATAT AAGCACTAAC ATTCCAACGT GCACATATTG GAACGACATC CATAACTCCA	8700
GAGAATCTCT AAAGTTTACA ATTTAAATGA ATTAACAATT TTCCCAACTA AAAGCACTCC	8760
AGTTACCGCA ACGATTTGTA CTGAATGTAC TAAATCGCAT TCCATCAACT TCATCTGTTT	8820
CGTCAACTTG AACAGACT AATTGAAGAT TTAATACTTC TTCTGCCATA GCTAGCTCCT	8880
CCTATTAAA TTTTGGGAT TAAGTACTTT ATCCACCCCTC ATTATACTCT CTCCACCACT	8940
AAAATGCAAG CAATTATACA ATGTTGTCAC ATAGAAAATA ATGTTTCCGT AACTTTCAA	9000
AGTAACCTCC ATCTCTCTCC CAAAACGTGA AGTTAGTTTT AGAAGTTACC TAAAAATCAG	9060
GTCACCTATT TPAAAAAGC AGCAAACTAT AAACTAGTAG GTTCCACACC AAATGTAGTC	9120
CCATACTGCC CCATAAGTCA GATTTATAGC GCACCATACC TAAAAACATC CCAAGTGAAA	9180
CATACAAACA CCAAGCTAGA ATGGTTCCTG TATGATGTGC TAAGGCAAAT AAAACACTTG	9240
TCAAAGCAAC TCTGATATCT AATTTTCTGA CCAAATTCCA TAAAATTTCT CGATACAGAA	9300
ATTCTCAAC CATAACTCGCA TTGATTAAGA ACAATAAAA TGAAAACCAA GGAATTTGAT	9360
GTTGAAGGCC AATTAAGTTT GTTGATTCG TGCTTCCTG ACCATGAATC AGACTAAAAC	9420
ATAGACTTAT AATCAGTAGG CTAACAAATT CAACACCAAG CCATTCATC CTAGATTCA	9480
TATTGACCTT ATGCCGTTGT TTGCGTTGGC CATACATCCA TAAAAAAGAA ATGAGTGACG	9540
AACCATAGAG AATCTGTAGT ATAGTTmACT CACCGATACA AAGAAATTTC AATAAGTATA	9600
GAGrTACCAA TASGACATTT ACTTGTGGG ATATATAAAC TGGAATTATT CTTTTCATAG	9660
TTACCTCCGA AATAATCTT CATAATCTAA ATCTAATACC TGCAACATCC TTT	9713

(2) INFORMATION FOR SEQ ID NO: 44:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 8657 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 44:

AAAGAAATTG TCAGAGAGTG GCTAGATGAA GTAGCAGAGC GGGCTAAGGA CTATCCAGAG	60
TGGGTGGATG TTTTCGAGCG TTGCTACACC GATACTTGG ACAATACGGT TGAAATCTTA	120
GAAGATGGTT CAACTTTGT CTTGACTGGG GATATTCCCTG CCATGTGGCT TCGAGATTG	180
ACAGCCCAAC TCAGACCCCTA CCTTCATGTA GCTAAAAGAG ATGCCCTCCT GCGTCAGACC	240
ATTGCAGGTT TGGTCAAACG TCAGATGACC TTGGTACTCA AGGATCCCTA TGCTAATCTCC	300

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TTCAACATTG AGGAGAACTG GAAAGGGCAC CACGAGACTG ACCACACAGA CCTTAACGGC	360
TGGATCTGGG AGCGCAAGTA TGAGGTGGAT TCGCTTGCT ATCCCTTGCA GTTGGCTTAT	420
CTCCTCTGGA AAGAGACTGG CGAGACTAGT CAGTTGATG AGATTTTGT CGCAGCGACT	480
AAGGAAATTC TCCATCTGTG GACGGTGGAA CAAGACCACA AGAACTCTCC TTATCGTTT	540
GTCCGAGATA CGGACCGTAA GGAAGACACC TTGGTAAATG ATGGCTTG ACCTGACTTT	600
GCAGTGACAG GTATGACTTG GTCAAGCTTT CGTCCGAGTG ATGACTGTTG CCAGTATAGT	660
TACTTGATTC CGTCAAATAT GTTGTGCTGA GTAGTCTTGG GTTATGTGCA AGAAATCTTC	720
GCAGCATTAA ACCTAGCTGA TAGCCAGAGT GTTATTGCTG ATGCCAAGCG TCTTCAGGAT	780
GAAATCCAAG AAGGAATCAA AAACATACGCT TACACCACCA ACAGCAAGGG CGAAAAGATT	840
TACGCTTTG AAGTGGATGG CCTAGGAAAT GCCAGCATCA TGGATGATCC AAATGTACCA	900
AGTCTACTAG CTGCGCCCTA TCTGGGCTAC TGTTCGTGC ATGATGAAGT GTATCAAGCT	960
ACTCGTCGTA CCATTTGAG CTCTGAAAAT CCATACTTCT ACCAAGGAGA ATACGCAAGC	1020
GGTCTCGGCA GTTCTCATAC CTTCTATCGC TATATCTGGC CAATCGCCCT TTCTATCCAA	1080
GGCTTGACAA CAAGAGATAA GGCAAGAGAAA AAATTCTTGC TGGATCAGCT GGTTGCCTGC	1140
GATGGTGGTA CAGGTGTCAT GCACGAAAGC TTTCATGTAG ATGATCCGAC CCTCTACTCT	1200
CGTGAATGGT TCTCCTGGC TAACATGATG TTCTGTGAGT TGGTCTTGGG TTACTTGGAT	1260
ATTCGCTAAG GGGCTCGCTT TAGCTCAACC GATTCTTATC AGAACACAAA GTTTACATTT	1320
AAAACGTTAA AATTAAATT TAGAATGAGG TTTTACTTCA TGGAAAATGT TGGTGTACAT	1380
ATTATCTCAC ATAGTCACTG GGATCGTGAG TGTTACTTGC CTTTGAAAG CCATCGTATG	1440
CAGTTGGTGG AATTGTTGA CAATCTCTT GATCTCTTG AAAATGACCC TGAGTTCAAG	1500
AGTTTCCACT TGGATGGACA AACTATTGTC CTTGATGACT ACTTACAAAT TCGCCCTGAA	1560
AATCGCGACA AGGTCCAACG CTACATTGAC GAGGGCAAAC TAAAAATTGG TCCCTTTTAC	1620
ATCTTGCAGG ATGACTACTT GATCTCCAGT GAAGCCAATG TCCGCAATAC CTTGATTGGT	1680
CAACAAGAAG CTGCCAATG GGGTAAATCA ACCCAGATTG GCTACTTTCC AGATACCTTT	1740
GGAAATATGG GACAAGCGCC TCAAATTCTT CAAAATCAG GCATTCACGT GGCGGCCTTT	1800
GGTCGTGGTG TGAAGCCGAT TGGATTGAC AACCAAGTCC TTGAAGATGA GCAGTTACG	1860
TCTCAGTTT CAGAAATGTA CTGGCAGGCT GTGGATGCTA GTCGTGTGTT AGGTATTCTC	1920
TTTGGCAACT GGTACAGTAA CGGGAATGAA ATTCCAGTTG ACAAAAGATGA GGCCCTGACC	1980
TTCTGGAAAC AAAAATTGTC AGATGTGCGT GCCTACGCTT CGACCAACCA ATGGTTGATG	2040
ATGAACGGCT GTGACCACCA GCCTGTACAG AAAAATCTGA GCGAAGCCAT TCGTGTGGCA	2100

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AATGAACTCT	TCCCCGATGT	AATCTTGTT	CATAGTTCTT	TTGATGAATA	TGTTCAAGCT	2160
GTAGAAGGTG	CGCTTCCTGA	ACACTTATCA	ACTGTTACAG	GCGAGTTGAC	CAGTCAGGAA	2220
ACAGATGGCT	GGTACACACT	TGCCAACACT	TCTTCATCCC	GCATTTACCT	AAAACAAGCC	2280
TTCCAAGAAA	ATAGCAACCT	CCTAGAGCAA	GTGGTAGAAC	CCTTGACTAT	TATCACTGGT	2340
GGACACAAACC	ACAAGGACCA	GTTGACCTAT	GCTTGGAAAA	CACTTTGCA	GAATGCCCA	2400
CATGATAGTA	TCTGTGGCTG	TAGCGTGGAC	GAAGTTCACCC	GCGAGATGGA	AACCGTTTT	2460
GCCAAGGTCA	ACCAACTAGG	AAACTTTGTT	AAAAGTAACT	TGCTAACGA	GTGGAAGGGT	2520
AAAATTGCTA	CGGATAAGGC	TCAAAGTGAC	TATCTCTTAA	CTGTCATTAA	CACAGGCTTG	2580
CATGATAAGG	TCGATRACTGT	CAGCACAGTG	ATTGATGTGG	CGACTTGTGA	TTTCAAGGAA	2640
TTGCACCCAA	CAGAAGGCTA	CAAAAAGATG	GCTGCTCTTA	TCTTGCAAG	TTACCGTGTG	2700
GAGGACTTGG	ATGGTCGTCC	TGTAGAGGCT	ACAATCGAAG	ACCTCGGAGC	TAATTTGAG	2760
TATAATTTCAC	CAAAGACAA	GTTCGGCCAA	GCTCGTATTG	CTCGTCAAGT	GCGCGTGAAC	2820
ATTCCAGTTC	ACCTAACGCC	GCTTCTTGG	ACAACCTTCC	AATTGCTGGA	AGGAAAACAA	2880
GAACACCGTG	AGGGTATTAA	CCAAAACGGA	GTGATTGATA	CACCATCGT	AACGGTGAGT	2940
GTGGATGACA	ACATCACAGT	CTATGACAAG	ACAACCTCACG	AAGCCTATGA	AGACTTTATC	3000
CGCTTTGAAG	ACCGTGGGAA	CATCGGAAAC	GAGTATATCT	ATTTCCAACC	AAAAGGAACA	3060
GAGCCAATCT	TTGCAAGCT	TAAGGGCCAC	GAGGTCTTGG	AAAACACAGC	TTGCTATGCT	3120
AAAATCTTGC	TCAAACATGA	ATTGACCGTG	CCTGTCAGTG	CGGATGAAAA	GCTAGAAGAA	3180
GAGCAACAAG	GTATCATCGA	GTTTATGAAG	CGTGAGGCTG	GACGGTCAGA	AGAATTGACA	3240
AAACATTCTC	TGGAAACTGA	GTGACTGTC	TTCGTTGACA	ATCCACAAAT	CCGCTTCAAG	3300
ACTCGTTTA	CTAACACTGC	CAAGGATCAC	CGTATCCGTC	TCTTGGTCAA	GACTCATAAC	3360
ACGCGTCCAA	GCAATGATTC	TGAAAAGTATC	TATGAGGTGG	TGACACGACC	AAACAAACCA	3420
GCTGCTTCAT	GGGAAAACCC	TGAAAATCCT	CAACACCAAC	AAGCTTTGT	CAGTCTGTAT	3480
GACGATGAAA	AAGGGTGCAC	TGTATCCAAC	AAGGGATTGA	ATGAATACGA	AATCCTTGGG	3540
GATAACACCA	TTGCCGTGAC	CATTTTGCGT	GCATCAGGTG	AGCTAGGTGA	CTGGGGCTAC	3600
TTCCCAACGC	CAGAAGCACA	ATGCTTGCGG	GAGTTGAAAG	TCGAGTTGCA	ACTTGAATGC	3660
CACCAAGCCC	AAGAACGCTT	CTCAGCCTAT	CGTCGTGCCA	AAGCCTTGCA	GACACCGTTT	3720
ACCAGCCTTC	AGCTTGCTAG	ACAGGAAGGA	AGCGTGGTTG	CGACTGGTAG	CCTCTTGAGC	3780
CATTCTGTT	TCAGCATACC	GCAAGTTGT	CCACAGCCT	TTAAGGTAGC	TGAAAATCAA	3840

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GAAGGCTATG TGCTTCGTTA CTACAATATG TGTAGTGAAA ATGTACGTGT GCCAGAAAGT	3900
CAACATCTCT TCCTTGACCT ACTTGAACGA CCATACCCAG TTCATTCAAGG ACTATTGGCT	3960
CCACAAGAGA TTCTGACAGA ATTCAATCAA AAAGAAGAAA TTAAATTCA AAAAGTAAC	4020
ATCAAAAGAA AGGAGGGCG AAAAGTAAG AACTAAGTGC TGATTCGCC CTTTTATGGT	4080
AAAAACAAATG ACCATTGCAA CGATTGATAT CGGAGGGACT GGGATTAAGT TTGCCAGTCT	4140
GACTCCTGAT GGGAAAATAC TGGATAAGAC AAGTATTCA ACCTTGAA ACTTGGAGGA	4200
TTTACTAGCG TGGCTAGATC AACGCTTGTC AGAACAGGAT TACAGTGGGA TTGCTATGAG	4260
CGTTTCAGGT GCAGTCATC AAGAGACAGG TGTGATTGAT GGCTTCAGTG CGGTGCCCTA	4320
CATCCATGGC TTTTCTTGGT ATGAGGCCT TAGCTCTTAT CAGCTACCTG TCCATTAGA	4380
AAATGATGCC AACTGCGTTG GACTCAGTGA ACTACTAGCT CATCCAGAGC TTGAAAATGC	4440
AGCCTGTGTC GTGATTGGG CAGGGATTGG CGGAGCCATG ATTATCAATG GTAGACTTCA	4500
TCGAGGTCGC CACGGCTCTGG GTGGAGAATT TGGCTACATG ACAACCCCTG CCCCTGCTGA	4560
AAAACCTTAAT AACTGGTCGC AACTAGCATC AACTGGGAAT ATGGTACGAT ACGTGATTGA	4620
AAAATCTGGT CATACTGATT GGGACGGTCG CAAGATTTAC CAAGAGGCC CAGCTGGTAA	4680
TATCCTTTGT CAAGAAGCCA TTGAGCGCAT GAACCGCAAT CTGGCGCAAG GCTTGCTCAA	4740
TATCCAGTAT CTGATCGATC CAGGTGTATC CAGTCTGGGT GGCTCTATCA GTCAAAATCC	4800
AGATTTTATC CAAGGTGTCA AGAAGGCTGT TGAAGACTTT GTGGATCCCT ACCAACAAATA	4860
CACGGTCGCA CCAGTTATCC AGGCCTGCAC CTATCACGCA GATGCCAAC TCTACGGTGC	4920
TCTTGTCAAC TGGCTACAGG AGGAAAAGCA ATGGTAAGAT TTACAGGACT TAGTCTCAA	4980
CAAACGCAAG CTATTGAGGT TTAAAGGT CACATTCTC TACCAGATGT GGAAGTGGCT	5040
GTCACTCAGT CTGACCAAGC ATCTATCTCT ATCGAGGGTG AGGAAGGTCA CTATCAATTG	5100
ACCTACCGCA AACCTCACCA ACTTTATCGT GCCTTGTCCT TGTTGGTAAC AGTTCTAGCA	5160
GAAGCTGATA AAGTAGAGAT TGAGGAACAA GCAGCTTACG AAGATTTGGC TTACATGGTT	5220
GACTGTTCTC GAAATCGGGT GCTGAATGTG GCTTCTGCCA AGCAGATGAT TGAGATATTG	5280
GCTCTCATGG GCTACTCAAC CTTTGAGCTT TACATGGAA ACACCTACCA GATTGAAGGG	5340
CAGCCTTACT TTGGCTATTT CCGTGGAGCT TATTCAAGCAG AGGAGTTGCA GGAAATCGAA	5400
GCCTATGCCA AACAGTTGA CGTGACCTTT GTACCATGCA TCCAGACCTT GGCCCACCTG	5460
TCGGGCTTGT TCAAATGGGG TGTCAAGGAA GTGCAGGAGC TCCGTGATGT AGAGGACATT	5520
CTTCTCATTG GCGAAGAAA CGTTTATGAC TTGATTGATG GCATGTTGCA CACGTTGTCT	5580
AAACTGAAGA CTCGCAAGGT CAATATCGGG ATGGACGAAG CCCACTTGGT TGGTTGGGA	5640

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CGCTACCTGA TTCTGAACGG TGTGTGGAT CGTAGTCTCC TCATGTGCCA ACACTTGGAG	5700
CGCGTGTGG ATATTGCTGA CAAATATGGT TTCCACTGCC AGATGTGGAG TGATATGTTTC	5760
TTCAAACTCA TGTCAGCGGA TGGCCAGTAC GACCGTGATG TGGAAATTCC AGAGGAAACT	5820
CGTGTCTACC TAGACCGTCT CAAAGACCGT GTGACTCTGG TTTACTGGGA TTATTATCAG	5880
GATAGCGAGG AAAAATACAA CCGTAATTTC CGCAATCATC ACAAGATTAG CCATGACCTT	5940
GCATTTGCAG GGGGAGCTTG GGAGTGGATT GGCTTTACAC CTCACAACCA TTTTAGCCGT	6000
CTAGTGGCTA TCGAGGCTAA TAAAGCCTGC CGTGCCAATC AGATTAAGA AGTCATCGTA	6060
ACGGGTTGGG GAGACAATGG TGGTGAAACT GCCCAGTTCT CTATCCTACC AAGCTTGCAA	6120
ATCTGGCAG AACTCAGCTA TCGCAATGAC CTAGATGGTT TGTCTGCGCA CTTCAAGACC	6180
AATACTGGTC TAACCGTTGA GGAGTTTATG CAGATTGACCC TTGCCAACCT CTTACCAGAC	6240
CTACCAGGCA ATCTCAGCGG TATCAATCCC AACCGCTATG TTTTTTATCA GGATATTCTT	6300
TGTCCGATTC TTGATCAACA CATGACACCT GAACAGGACA AACCGCACTT CGCTCAGGCT	6360
GCTGAGACGC TTGCTAACAT TAAAGAAAAA GCTGGAAACT ATGCCCTATCT CTTTGAAACT	6420
CAGGCCAGT TGAATGCTAT TTTAAGTAGCC AAAGTAGATG TGGGACGACG CATTGTCAG	6480
GCCTACCAAG CGGATGATAA AGAAAGTTA CAACAAATCG CCAGACAAGA ATTACCAGAA	6540
CTTAGAAGCC AAATTGAAGA CTTCCATGCC CTCTTTAGCC ACCAATGGCT GAAAGAAAAC	6600
AAGGTCTTTG GTTTGGATAC AGTTGACATC CGTATGGCG GACTCTTGCA ACGCATCAA	6660
CGAGCAGAAA GCGGTATCGA GGTGTTATCTG GCTGGTCAGC TTGACCGCAT CGACGAGCTG	6720
GAAGTTGAAA TCCTACCATT TACTGACTTC TACGCAGACA AGGATTTCGC AGCAACTACA	6780
GCCAAACCGT GGCATACCAT TGGCACAGCG TCGACGATTT ATACGACTTA ATATTCTTCG	6840
AAAATCTCTT CAAACCACGT CAGCTTCCAT CTGCAACCTC AAAACAGTGT TTTGAGCAAC	6900
CTGCAGCTAG CTPCCTAGTT TGCTCTTGA TTTTCATTGA GTATAAAAAC AAGAACACCT	6960
TGCTTGGCGC AGGGTGTTC GCCTGAAACA GAAGAATTAT CTGGTTCAA ATGCTACAGT	7020
TAGACAAACT TATGATAAAA TAGCAGAAAG TGAATGTTTC CTAAGAGCAA TTGGAGGTAT	7080
TATGCTACAC TTAAAATTAG TAAACACAAGA AATAGAAGCT GAAAAGCCAG CATCTGTAGA	7140
AGCTTGGATC ATTTCCGTCA AATTAAAAAA AGGTGCTAC CGACATATAT AGATTCCAAA	7200
AACAAAAACG TTACCGGAAC TAGCAGATGT GATTTTATGG AGTTTGATT TTGCAAATGA	7260
TCATGCTCAC GCATTTTCA TGGATAATGT TGAGTGGAGT CATGCAGATT CTTACTTCG	7320
TAGCTTGTGTT AGTGACGATG TTGAAGAACG TTACACAGAA AATGTCTATC TGGATAGCCT	7380

AAGTGTCAAA CAAAAATTAA AGTTTATTT CGACTTCGGT GATGAATGGC GTTTGAAATG	416 7440
CCAAGTGCTG AGAGAAATCG AGACAGAGGA CGAAGAAGCT TATCTCGTAC GTTCGGTTGG	7500
AACGTCGCCA GAACAATATC CAGATTATGA TGTTTGTAC TATGAAGAAT GGAAAATTG	7560
AAATCAGTCT GTGTAGGCTT AGTATTCAA TAGACTTCCT GCACAACTAG AATCCTAGTT	7620
CATGATTGAT AATACCAGCA ATCAAATTCA TTCGTAATCC GAAGCGTTA CGATGATTTC	7680
GATAGTTGT TGAAAACATT TTAAACGTTT TTACTTTGGC AAAGATGTTC TCAACCTTGC	7740
TTCTCTCCTT AGATAGCGCA TGGTTATAGG CTTTATCTTC AGCTGTTAGT GGCTTGAGTT	7800
TGCTGGATT ACGTGAAGTT TGTGCTTGAG GACATATCTT CATGAGCCCT TGATAACCAC	7860
TGTCAGCCAA GATTTACCA GCTTGTCCGA TATTCCTGCA ACTCATTTG AACAACTTCA	7920
TATCATGACA ATAGTTCACCA GTGATATCCA AAGAAAACAAT TCTCCCTGAA CTTGTGACAA	7980
TCGCTTGAGC CTTCATAGCG TGAAAATTCT TTTTACCAAGA ATCATTGCT AATTCTTTT	8040
TTAGGGCGAT TGATTTTAC TTCCGTCGCA TCAATCATTAA CCCTGTCCTC AGAACTAAGA	8100
GGAGTTCTTG AAATCGTAAC ACCACTTTGA ACAAGAGTTA CTTCAACCCA TTGGCTCCGA	8160
CGGATTAAGT TGCTTCGTG AATACCAAAA TCAGCCGCAA TTTCTTCATA AGTGGGTAT	8220
TCTAGGCTTA ATTAGGTTT TCGTCCACCT TTTGCGTGTGTT TAAGTTGATA AGCTGTTTTT	8280
AATACAGCTA ACATCTCTTT AAAAGTCGTG CGCTGAACAC CAACAAGACG CTTAAATCGT	8340
GTATCAGTTA ATTGTTTACT TGCTTCATAA TTTCGCAGGG AGCTTATTGA CTCTTTGGTA	8400
GGTGTCAATG TTTTTTCAT CTATCCCGAG AATTATTTTC CCGCCATTG TATTGCAAA	8460
TGCTGAGTAG GTTTCCAGA AAGACTCTGG AAGATTGTTT TTAGCTTTTT TGTATTCTAA	8520
ATCAACCCCT TCAAATTAA AGTCCATATT TTTCTTTAC ATCTGTTTT TGTGGTTCTG	8580
GTATTGTTCAAGTGAGTG ATAATATAGC GAATTGAATT TCGAGAGTTT TTACTCAGTT	8640
AATTTCTTTT TTAACCC	8657

(2) INFORMATION FOR SEQ ID NO: 45:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 11384 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 45:

TCTATTTGG GTATAGACTT ACCTATAAG AAAAATATCT ATACACTGCC TTACTAGCTA	60
TACTGAACGA GTCAACAAAA ACGATATATA TTGATGATAT AAATACAGCA AGATTTTTA	120

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ACTTCTTGG CAATGATATT CCTAATTCGT CTTAAAAAA AATTGACTAT ATCGCACCTT	180
CAGAAATTGT TTCATTTAGT ACCTACGTTC GACAACGTTC TAAAGTAATT CCTAAAATT	240
TGGAACATAT ATTAAAATCA AGTTTTTAT TAGAGAATAT AGATGTTCT GGTTACACTG	300
TAAATATTT AGAAGATCAA TTAACAAAAC ATAGAACATA CAAAATTAGT AAAACTAAC	360
TGGTTGATCT CATGTATAAA TACCTAACAA AACCACGCC CTTGCCGTGCT GATGGAAAGA	420
AAGGTACAAA TACATGAATA TCAAAAGAAAA AATCAAAAG AATGGCCAA GAGTTTATTA	480
TGCTAGTGTT TATCTAGGCG TTGACCAACT AACGGGCAAA AAAGCCCGTA CAACTGTTAC	540
ACCAACCACT AAAAAGGGCG TTAAAGTAA AGCGCGTGT GCGATCAATA CTTTGCTGC	600
TAATGGCTAT ACAGTTAAAG ACAAGCCGAC AATTACAACA TATAATGAGC TTGTTAAAGT	660
TTGGTGGGAT AGTTACAAGA ATACAGTTAA GCCAATACT CGCCAATCCA TGGAGGGATT	720
GGTTAGAGTG CATTATTGC CTGTATTTGG CGATTACAAG CTATCTAAC TTACTACGCC	780
TATTCTTCAA CAGCAAGTAA ACAATGGGC TGACAAGGCA AATAAAGGCG AAAAAGGGC	840
ATTTGCTAAC TACTCTTGC TCCATAACAT GAATAAGCCT ATTTTGAAAT ATGGCGTAGC	900
TATCCAGGTA ATACAATACA ACCCAGCTAA TGATGTCATC GTTCCACGCA AACAGCAAA	960
AGAAAAGGCT GCTGTCAAAT ACTTAGACAA CAAAGAATTA AAACAGTTTC TTGATTATTT	1020
AGATGCTCG GATCAATCAA ATTATGAGAA CTTATTTGAT GTTGTCTGT ATAAGACTTT	1080
ATTGGCCACT GGTTGCCGTA TTAGTGAGGC TCTGGCTCTT GAATGGCTG ATATTGACCT	1140
AGAAAGCGGT GTTATCAGCA TCAATAAGAC ACTAAACCGC TATCAGGAA TAAACTCACC	1200
TAAATCAAGC GCTGGTTATC GTGATATACC AATAGACAAA GCCACATTAC TTTTACTGAA	1260
ACAATACAAA AACCGTCAAC AAATTCAGTC TTGGAAATTA GGCGATCTG AACAGTTGT	1320
ATTCTCTGTA TTTACGGAGA AATATGCTTA TGCTTGTAAC TTACGCAAC GCCTAAATAA	1380
GCATTGAT GCTGCTGGAG TAACAACTGT ATCATTTCAT GGTTCCGCC ATACACATAC	1440
TACTATGATG CTCTATGCTC AGGTTAGCCC GAAAGATGTT CAGTATAGAT TAGGCCACTC	1500
TAATTGATG ATCACTGAAA ATACTTACTG GCATACTAAC CAAGAGAATG CAAAAAAAGC	1560
CGTCTCAAAT TATGAAACAG CTATCAACAA TTTATAAAA ATAAGGGTGA CCCATTTCCG	1620
GGCTACCCCTC TTACTATACC AAAAATTAGT AGGGGTAGTA AAAAGGGTAT TAAATTATAA	1680
AAAGCACTAA GGGAAAGGCC CCCAAAGTGC TTATTCAAA GGCTTTATAG CCTATAATCA	1740
CATAAAAGAGA TTATTTTTA AGGTTGTAGA ATGATTCAA TCCACGATAT TCAGCTACTT	1800
CACCAAGTTG GTCTTCGATA CGAAGCAATT GGTTGTATTT AGCGATGCCG TCTGTACGTG	1860

418	
AAAGTGAACC AGTCTTGATT TGTCCTGCGT TAGTTGCAAC TGCAATATCA GCGATTGTTG	1920
AATCTTCAGT TTCACACTGAA CGGTGTGATA CAACAGCAGT GTAACCAGCT TCTTTAGCCA	1980
TTTCGATAGC TTCAAAAAGTT TCAGTAAGAG TACCGATTTG GTTAACCTTG ATAAGGATTG	2040
AGTTAGCAGC ACCTTCTTGG ATACCACGTG CAAGGTAGTC AGTGTGTTGT ACGAAGAAGT	2100
CGTCACCAAC AAGTTGTAAC TTCTTACCAA GACGTTCACT AAGAGCTTTC CAACCATCCC	2160
AGTCGTTTC ATCCATACCA TCTTCAATAG TGATGATTGG GTATTTGTTA ACCAATTCTT	2220
CAAGGTAGTC GATTGTTCT GCAGATGTAC GAACAGCAGC ACCTTCACCT TCAAATTTAG	2280
TGTTAGTCGA AACTTTACGT TCTTTATCGT AGAATTCTGA TGAAGCACAG TCAAATCCGA	2340
TAAATACTGC TTTACCTGGT ACATATCCAG CAGCTTCAAT CGCAGCAAGG ATAGTTTCAA	2400
CACCATCTTC AGTTCCCTCG AAACGAGGAG CGAATCCACC TTCGTCACCT ACGGCAGTTT	2460
CCAAACACG TGATTTAAGG ATTTCTTAA GAGCGTGGAA GATTTCAGCA CCGAACGAA	2520
GGGCTCTTT AAATGTTGGC GCACCAACTG GCAAGATCAT GAACCTCTTG AAACCGATTG	2580
GAGCGTCAGA GTGAGAACCA CCCTTGATGA TGTCATCAT TGGAGTTGGA AGAACTTTAG	2640
TGTTGAATCC ACCAAGATAG CTGTAAGTG GGATTTCAAG GTAGTCAGCA GCAGCACGAG	2700
CTACAGCGAT AGACACACCG AGGATTGCAT TCCGACCCAA TTTACCTTGT TTAGGAGTAC	2760
CGTCAAGTGC GATCATAGCA CGGTCAATAG CTTGTTGATC ACGTACATCG TAGCCAATGA	2820
TAGCTCAGC AATGATCTTC TTACCTTOT CAACAGCTTT TTGTGTACCA AGACCACCGT	2880
AACGAGATTG GTCACCGTCG CGAAGTTCAA CTGCTTCGTG TTCACCAAGTA GAAGCTCCTG	2940
ATGGAACCAT ACCACGTCCG AAACGACCTG ATTCACTGTA AACTTCTACT TCAAGTGTG	3000
GGTTACCCCG TGAGTCTAGG ACTTCGCGAG CGTAAACATC AGTAAATAATT GACATTTTTT	3060
ACTCTCCTTA TGAGTTAAAT TTTTACACC TCTATAATAC CTTAAACCC CTCCTTTTC	3120
AAGAAAAAAC GTTATCTTGT TGCAACTTTT CCTTAACCTTT ATAAAGTAAT CGCTTCTTT	3180
TGTCTGTTT ATTCTAACTT TTATGATATA CTGTTTCTAT GACAGATTAA TCAAACAAAT	3240
TACTTGAAAA AGCTCATGGT GGTTAAAAAA TAAATCCGGA TGAGCAAAGA CGCTATCTTG	3300
GTACTTTGA GGAAAGAGTT CTTGGATATG TAGATATTGA CACAGCAAAT AGCCCTCACT	3360
TAGAAAAAGG CTTTTTATTT ATTTTAGAAA ACCTTCAGGA AAAAGCAGAG CCACTATTG	3420
TGAAGATTTC ACCAACTATC GAAMTGATA AGCAAGTTT CTACTAAAA GAAGCAAAG	3480
AAACTGATAG TCAAGCCACC ATAGTATCTG AAGAGCATAT TACTTCTCCT TTTGGCCTGG	3540
TTATTCTATAG CAATGCACCA GTTCAAGTAG AAGAAAAAGA CCTTCGACTT GCTTTCCAA	3600
AACTTTGGGA AGTTAAAAG GAAGAACAG CAAACACATC CTTATGGAAG AAATGGTTA	3660

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GCTAAATCTT GCACATATTT AATAAGTGCC CAATATTGGC AGCCGTGCGC TCCAGATAGA	3720
AACTGGCATT TTTCAAACTA TCTTCTAAAG GTTCACTTTT CTCCAAAATA GAAAAGACAG	3780
CTTGGATATT TTCAAAATGGT AGGGGAGGTA AATCTTCAGC AAGACTACCG CAAATAGCAA	3840
TAACAGGAAC TCCAACAGGG GTTCTTTTG CAACACCTAT AGGCGCTTTC CCAGCAAAGC	3900
TTTGACTATC AAGTCTTCCT TCTCCAACAA CAACCAAGTC AGCATCTGAA ACTTTCTTAT	3960
CAAAGTTGAT TAAGTCCAAG CAGGTATCAA TTCCAGACAC GATACTGCC TGAGCAAAGG	4020
CACACAAACC ACCAGCAAGG CCTCCACCTG CTCCTGCTCC TTTAATTCT AATGTTGCAG	4080
GTGAGAATTT TTCATAAAAA TCTTGGATCG CCTGATCTAC GACTGCAAAC ATAGTCGGAT	4140
GTAGACCTTT TTGATTGCCA AAAGTGTAAAG TCGCACCTG ATGACCACAT AAGGGACTCA	4200
CGACATCTGC TAAAATATGA ATTTGAACAC CTTCAGGAAT TTTATAGCAA TTTCTGTTG	4260
AAACAGAACG TAAGTTTAAT AAGGATTGAC CGGAAGCAGG CAAGACATT CCATCCCTAT	4320
CATAAAATTG ATAACCTAAA CCAGCAGCAA TCCCCAGTCC TCCATCATTA CTGGCCGTGC	4380
CACCAACACC GATATAAATA TCTTAAATCC CTTTAGAGAT GAGATGAAGA ATCAACTCTC	4440
CAATACCACA AGTTTGGATT TGAAGTGGAT TTCGTTCTC TAGCGGAATT TTTCCAAGAC	4500
CAACCAAGTC AGCTACTTCA AATAGTGCCA GTTCCCCTTT TTGAAAATAG CGCATGGCTT	4560
CTTTTTGTCC AAAAGGGTCT GTCACTTGGA TCCATTTTC TTTTAGGTCA AGAGAATGTC	4620
GGATAGCATC TACAGTACCT TCTCCCCAT CACCAACAGG GCAGAGGAGA CATTCTACAT	4680
CTGCTATCGA TTGTTGGAAG CCTCTTTTA TTGCTTCAGC TACCTGTTGA GCTGTCAAGC	4740
TTTCCTTAAA CGAATCCGGT GCAATTACAA TCTTCATATT TTCCCTCATT CTAACAGTC	4800
AATCAAAGGG AGAACTCTA AAAAATCCCT CTTGTCAACA TGATGTGGTA TTTCTTTTTT	4860
GAGCACTTCT TTGGCACAAA AGGCGATTCC TAACTTCGCC GACTTCACA TTAATAGATT	4920
ATTAACCCCA TCACCGATTG CCACCGTTCT TTCTTTAGAA AGTTTTAGTT TCTTCTCCA	4980
TTTTTCCAGA GTCTCTTTT TGACCTGGGG ACTTATAATT TGTCCAACTA ATTTCTGT	5040
TAAAAGACCT TCTTGACTT CAAGCTAGTT GGCAGTGAAA TAGGCAATAC CAAGGGATT	5100
TGCTAATCTC TCCAACATT GGTGTAAATC CACCAGACAC CAGACCAACT AGGATGCCAT	5160
TCTTTGGAG AATAGAGATG AACTCTGGGA CATTAGCGA TAGATGAATT GAGTTGAAGA	5220
CGTTATCAAA GACCAAAATA GGAAGACCTT CCAACAAAGGA CACTCTTTT CTTAAACTGC	5280
TTTCAAAGAC CAACTCTCCT CGCATTGCTC GACTTGTAAAT CTGCGAAATT TCCGCCTCAT	5340
GACCTGCCTC TCTCCCTAAA AGATCAATCA CTTCTCTAG GATTAAGGTT CCATCTACAT	5400

420	
CCAAAACACA CAAGCCTTT ACTTGAGACA TCAGTCTCC TCTCTAAACA GCCTAAAAAT	5460
CGTATGAAGT CATCATACGA TTTTATCTAT TAATTAACTA AACTATGGTA CAAGTCAGG	5520
TATGACTTGC AGGCTGTATC CCATGAGAAG TCACTCTCCA TAGCTGTTT TTGTAGGTTT	5580
CTCCAAATGT CTGGATGGTT TCTATAACAAG TCCAATGCTG TTTGGAAAGT CCAATTAAAC	5640
CAATAAGGAG ATAGATTGTC AAAGCTAAAG CCAGTACCGC TTCCCTCGAT TGGATTGAAA	5700
GCGCGAACTG TATCTCGCAA GCCTCCAACT TCATGGACCA ATGGCAAGGT TCCATAACGC	5760
ATAGCCATCA TTTGAGACAA CCCACACGGT TCAAAACGAC TTGGCATGAG GAAGAGGTCA	5820
CAAGCAGCGT AGATTCTCTG AGCAAGTTG ACATCAAAG TGATATTGTT TGATAGCTTG	5880
TCTGGTAAA TCTGAGCAAA CCATGAGAAA CCTCCTCAA AGGCTGGATC GCCAGTTCCC	5940
AAAAGAACAA TCTGAACATC TTCTTGCAG ATATGGTCAA GACTTTGAC CACCACATCA	6000
AAACCTTTT GACGTGCAA ACGAGAACAA ATCCCCACCA GTGGAACGTC TGCTCTAACCA	6060
GGCAAGCCAA CTCTTCTTG CAATTTGCC TTATTTTGG CTTTCCCAGA CAAATCTTCC	6120
TGATTGAAAT GATAGCTAA AAGAGCATCC GTCTGAGGAT TATAAAGATC AGCATCAATC	6180
CCATTACGA TACCAGATACT TTTACCAGAC TCCATTAA GAATCTGATC CAAATTACAT	6240
CCAAACTGAC TAGTCATAAT TTCATGAGCA TAGCTAGGTG AAACGGTTGA AACACGGTTC	6300
GCATAGAGAA TACCTGCCTT CATCCAGTT AGACAGTTG TCCATCGAAG GGTGCCATCA	6360
GCGTAACGTT CAAAGCCAAAC TCCAAACAAA TCACCCAAAC TTGCTTCTGA AAAATTGCTCT	6420
TGGAATTCTA ATTATGAAT GTTAAACT GTTCAATGT CCTCATAGGC TTGAATCCAA	6480
CGGTATTTT CCTCAACAA GAAAGGAATC ATAGCTGTAT GGTAGTCATG AACATGGAGA	6540
AGATCAGGAA TAAAGTCAAT CCTTCCATA GCCTCAATGG CAGCCAGTTG GAAAAGGCA	6600
AAGCGTTCTC CGTCATCAAA ATCACCGTAA ACATGACCAC GGAAGAAATA ATATTGATTG	6660
TCAATAAGT AGAAGGTTAC ACCATTAAAT ACTGTTTCT TAATTCCACA ATACTGTCTG	6720
CGCCAACCAA CGCTCACCTC AAAATGAAGC ACATCTTCAA TCTGATTCC AAATTTAGCC	6780
TCTACCATAT CATAGTAGGG TAAAATCACT GCAACTTCGT GCCCAGCTTT TACCAAGTGAT	6840
TTTGGAAAGAG CGCCAATGAC GTCTCCAAA CCACCTGTTT TTGAAAAGGG TGCAACCTCT	6900
GCTGCTACAA ATAAAATTTT CATGAATGAA TATCCTCTGT TACTTTAGCA CCTTTCTTAA	6960
CCACAACCTGG ATGTTCTGCA GTTCTCGAA TCACAAACACC ATGCTCAACT TCAACCCCTT	7020
TGTCCAAGAT AGCATATTCG ACCTGAGGCC CTTCTCCAAT AACACACGA GGGAGAGCA	7080
GGCTATCTTT AACCAAGCTA TCCTTATGGA CATGAATATT ACGTGATAGA ACAGAATTAG	7140
CTACTTGACC TTCAATAATA CTACCAGAGG CAAACTGAGA AGTGCCTTACCTTAGATGTAT	7200

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TAGCATAGTA AGTGGCTCT TCGTTTTGA CCTTTGTATA AATCTTTGG TTTGGTGAGA	7260
AAAGAGAATA GAATTTTGT GATTCAAGCA TATCGATATT CGCTTGATAA TAAGATTAA	7320
CAGAGTGAAT ATGGCTAGA TAGCCCGTGT ACTCGTAGGC GAAAGCTCCC TCTTTTACAG	7380
CCAAATCCCG TAAAACATAG CGCAATTCT CTGGATGTT TCCTTCTTCCA	7440
AGTGTCAAT CAACCAAGGT GTATCAACGA CAAAGATATC TGAGACATA TTGAACGTTT	7500
CAGCTGTTGA CTTGCTATCA AAGAGTTAT GAGAAAGAAC ATGGCTGTT TCATCTACAT	7560
CCAAGATTGC ATTACTTCT GAAATATCTT TCTTAGCTAG TTTTTTATAA ACTACAGTGA	7620
TAGGCTCTT TGTGACTA TGAGGTGGA AAACCTGGTT CAAATCAATG TTAATAAGAA	7680
CATCGCAGTT GAGGCCAAC GTTGGTTTG AGCCAGAACG TTTCAAATAA GTAAGAACG	7740
GTTGGTAGTA TTCTTTCCA ACTGTAACAGC TTTCTACACG GGTATTGTA ATTCCCTAGAT	7800
AGTAATGGCT AAGAAGGGTT GATAAGCCCC ACTCGCGTCC TGAACGAATA TGGTCAAATA	7860
CTGAGCTGAT ATTATCCTGC TGAAAATAC CAAAGACACT ACAGAACCT GCATTAGCAA	7920
GGCTTGAAAG TGGAAAGTCA ATCAAACGAT ATTTCACCAC AAATGGCAA CTTGCTACTG	7980
GACGGTGGTC CGTCATGTC GACATATTGT GAAAACCAAC TGTATTCCCT AAAATGGCAG	8040
AATATTATC AATCTTCATC TGTTGCTACC CCCACTACTT CATTATATCC TACAACTTGT	8100
ACTTCATCTG TTCCATCAAT TTCGACACCG TCAGAAATAA TCGCACCTTC ACCAATAATG	8160
GCACCTTAA TCTTAGCTCC TTGACCAATG ATAGCTCCAC TCATGATAAC TGAATCAAGG	8220
ACTTCCGCTC CTTCCGAAC TTGCGCCCT GTTGAAGGA TAGAATGTT AACAGTCCA	8280
TCAACGAAAC ATCCGCTAC AACTAATGAG TCTTCCACAT GAGCATTG CCGGAGGAAG	8340
TTTGGTGGTG AAATCAAGTT TCTTGAGTAA ATCTTCCATT GACGGTTACG ACTATCCAAG	8400
GCATTTCTG GAGAAATATA CTCCATGTT GCTTCCAAA GTGACTCAAT AGTACCAACA	8460
TCTTCCAAT ACCACTAAA TTCGTAAGCA TAAACACTTT CACCTGACTC AAGGTAATT	8520
GGAATGACAT TTTACAAA GTCTGACATG CCAACCTGC TCTTTTCAGC AGCGACTAAC	8580
ATATTACGAA GGCGMTGCCA ATCAAAATG TAGATCCCA TAGAAGCTT TGTAGATT	8640
GGTTGAGCTG GTTTTCTTC AAATTCAACA ATACGATTGT TAGCATCTGT GTTCATGATA	8700
CCAAAACGGC TTGCTCTTT AAGAGGGACG TCTAAAATG CTACTGTCAA CCTGGCATT	8760
TTATCCTTAT GAGACTGGAG CATATCATCA TAGTCCATT TGAGATGTT ATCCCCAGAC	8820
AAAATCAAGA CATACTCAGG ATTGACACTG TCGATATAGT CGATATTTG CTAAATAGCG	8880
TGACTAGTCC CCTCAAACCA ACGATTTCCCT TCACCTGAG AATAAGGTTG AAGAATAGAG	8940

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ACACCTGAAT TAATACCGTC TAGTCCCCAG CTTGAACCAT TCCCAATATG GTTGTGAGA	9000
GCAAGTGGTT GATACTGTGT AACGACCCCA ACATTGTGAA TCCCTGAGTT GGCACAGTTT	9060
GATAGGGCAA AGTCAATGAT ACGGTAGCGC CCACCAAATT GCACAGCTGG TTTTGCATG	9120
CTTTGAGTGA GTTTACCGAG ACGAGTTCCCT TGCCCACCAG CAAGAACATCA AGCTAACATT	9180
TCATTTTCA TTTTCTACTC CTTTTGGTT TTTATTTGTG ACGGTTTTAG TAGATTTCAA	9240
GCGACGTTG ATTTCCATA CACTTGCTCC CATAGCCGGT AGGGTAAAGG TTAAGGTCTG	9300
CTCATAATCT TTCCATAGTC CTTCTTGCCT TTGAACAGTT TGATTATGTT CTTTCCAAC	9360
GCCTCCCCAC TCTTCAACT CAGTATTCCA TACTTCTTCG TAAATTCCCTG CAACGGTAG	9420
TCCGATTGTA AAATCTTCC GCTCAACAGG TACCATATTA AAGATACAGA CTAACATTTC	9480
TCCCTTTTA CCCTTACGAA TAAAGGAAAG AACACTCTGG TCTCGATTAT CCGCATCAAT	9540
GATTTCAATA CCATCATAGC TGGAATCAAT TTCCACAGA CAGCGATGAT CTTTGTAAA	9600
CTGGTTTAGC TGAGAACGGA AATACTTCAT CTTAGCATTC ATTGGGTCTT CTAGGTTAGA	9660
CCATTCCAAC TGTCTTCAG ATTTCATTC TAGGAATTGA CCGTATTCGC TACCCATGAA	9720
GAGCAATTTC TTACCAAGGGT GACAAATTG GTACGTATAG AGATTGCGCA AGCCTGCGAA	9780
TTGATTGTA CGATCTCCCC ACATCTTATG CATCATACTC TTCTTGCCAT GAACCACTTC	9840
ATCGTGCAG AATGGCAAGA GATAATTCTC CTTGAAAACA TACATAAAAGC TGAAAGTCAC	9900
CAGGTTAAAG TCATATTTAC GATAGATCGG ATCTTCTCG TAGAAACGGA GGATATCATT	9960
CATCCAGCCC ATGTTCCATT TGTAGTCAAA TCCTAGACCA CCAATCTCTT TCATTCCCGT	10020
AATCTTGATC GCAGACGAAC TTCTTCTGC AATCATCATC ACATCTGGAT ATTCTAACCTT	10080
AATAACCTCA TTCAAGCGCT GAAGGAAATA ATAACCTTCATAG TGTAGTCAAA TCCTAGACCA CCAATCTCTT TCATTCCCGT	10140
TTTATTAGGT GTCCATGGAG CATCATCATA GTCCAAATAG AGCATGTTGC TAACAGCATC	10200
CACACGAATA CCATCCAAT GATAGACATC AATCCAATGC TTAATGCAAG AAATTAAGAA	10260
GGACTGGACT TCATTTTTC CAAGGTCAAA ATTAAGGGCA CCCAACCAT GGTTATGAGC	10320
CTTATTATGG TCTGGTATT CAAAAGTCGG TGTCCCATCA TAATAGGCTA AGGCATCATC	10380
GTTGATGGTA AAGTGACTGG TACCCAGTCC ACAATAACCC CAATATTATG GGTATGACAC	10440
TCCTCGACAA AATCTTGAAA CTCCCTCTGGT CGGCCATAAG CATGCTCTAA AGCGAAGTAA	10500
CCCATAAGCT GATAACCCCA ACTCAAGCCC AAAGGATGGG ACATCAAGGG CATAAAACTCA	10560
ATATGAGTAT AGTCATTTCA AACGAGATAA GGAATGAGTT CATCCTTGAG CTGGGCAAA	10620
CTATAAGGAC TGCCATCAGA ATTTCATTCAG CAGTGCAG CGTGAACCTTC ATAAATATTG	10680
ACAGGACGCT CTTCAAAGCC CCAACGTTTT CTTCGTGCCA GCCAAAGTCC ATCCTTCCAT	10740

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TTCTTCTCA	GAAGCTCTGT	TACGATTGCC	CCTGTTCTG	GACGAGCCTC	ATACCTGACA	10800		
GC	AAAAGG	GT	CAATCTTCAT	CAGTTGATGA	CCATTTGAC	GTGTGACATG	ATATTTGTAA	10860
ATATGCC	CTT	GAGCCAT	ATTGGTAAAG	ACTTCCCAGA	CCCCAAAATC	ATTTCTTACC	10920	
ATTGGA	ATCT	GATTTCAAT	CCAGTTGGTA	AAATCACCAA	CCAAGTGAAC	AGCCTGAGCA	10980	
TTAGGTGCC	AAACACGGAA	GGTATAGCCA	TGCTCTCCAT	TTAGTTCTTC	CCTATGTGCT	11040		
CCTAGATA	AT	GTTGAGATA	AAAATTTCA	CCCGTCATAA	AGGTTTTAA	TGCTTCTCTA	11100	
TTATCC	CATAT	ACTCCCCTTC	TCCTGTAAGC	TTTTCTATG	TTTTTATTAT	ACTACCTTT	11160	
TAGAGA	AGAT	TCAAGTAAT	TACTATACTT	CTTTAATTAT	TTTGAAAATC	TACAACAAGT	11220	
TCAC	TTACTC	GTTCAATTGT	AAATCAATAT	TTTTCAAAA	AATTGCGAAA	ACGCC	TTCT	11280
TTTTCTACTA	TAGT	GAAATG	AAATAAAACA	TGCGCAAATC	GATTAAGGAA	TTTAATCTAA	11340	
TTTCTAACAA	TGT	CTTAGAA	ATCAAAGTGT	ACTATTTAA	CTCC		11384	

(2) INFORMATION FOR SEQ ID NO: 46:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 7577 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 46:

TGTTGATTG	TTACTAGACG	TTGACCAACG	TCCTTCGGCT	GGAAAAGGAA	TTCTCCTTAG	60		
TTTCCAA	CAC	GTTTCGCCA	TGTTTGGTGC	GACCATCTG	GTACCATTGA	TTTGGAAT	120	
GCCTGT	ATCT	GTTGCC	TTT	TGCTTCAGG	TGTTGGAACA	CTCATCTACA	TGATTGCTAC	180
TGGTTTAA	AA	GTTCCAGTTT	ATCTAGGTT	TTCATTTGCC	TTTATCACAG	CTATGTC	ACT	240
GGCTATGAAA	GA	AAATGGGGG	GGGATGTATC	TGCTGCCAA	ACAGGGTTA	TCTTGACTGG	300	
TTTGGTCTAT	GTC	CTTGTTGTT	CTACCAGCAT	CCGATTGTA	GGAACAAAAT	GGATTGATAA	360	
ACTCTGCGA	CCAATCATTA	TCGGTCTAT	GATCATCGTT	ATCGGTCTTG	GACTTG	CAGG	420	
TTCAGCTGTT	ACCAATGCAG	GTCTTG	TAGC	AGACGGAAAT	TGGAAAATG	CTCTGGTAGC	480	
CGTTGTTACT	TTC	CTTAATTG	CTG	CTT	TAAT	AA	GGAAAAGGCT	540
CATTCCATTC	CTCTTGCC	TTATCGGTGG	TTAC	CTTTC	GC	ACTAACTC	TTGGCTTGGT	600
TGACTTTACA	CCAGTTCTTA	AAGCCA	ACTG	GTC	GA	AAATT	CCTGGTTCT	660
TAGCACAGGT	GGT	GCC	TTTA	AAGAGTACAA	TCTT	TACTTT	GGTCCAGAAG	720

424	
CTTGCCAATC GCTATCGTAA CAATTCTGA ACATATCGGA GACCATACTG TTTTGGGTCA	780
AATCTGTGGT CGTCAATTCT TAAAAGAACC AGGTCTTCAC CGTACTCTTC TTGGTGACGG	840
TATCGAACT TCTGTTCTG CCTTCCTGG TGACCAGCC AATACAACCTT ACGGAGAAAA	900
TACAGGGTT ATCGGTATGA CTCGTATCGC TTCTGTCTCA GTTATCCGTA ACGCTGCCTT	960
CATCGCGATT GCCCTCAGCT TCCTTGGTAA ATTCACTGCC TTGATTCAA CTATTCCAAA	1020
CGCTGTACTT GGTGGTATGT CAATCCTTCT CTATGGGGTT ATGCCAGCA ATGGTTTGAA	1080
AGTCTTGATT AAAGAACGTG TTGATTTCGC TCAAATCGA AACCTCATCA TCGCAAGTGC	1140
TATGTTGGTT CTTGGACTTG GAGGAGCTAT CCTTAAACTT GGTCCAGTTA CACTTCAGG	1200
TAATGCCCTT TCAGGCCATGA CAGGAATCAT CTTGAACCTTG ATCTTGCCAT ACGAAAATAA	1260
AGACTAAGAG TCTAAATACA CCTAATCCAC TCAGACAGCT GAGTGGATTT TTCGTATACC	1320
ATAATAAAAG TGTCTTAACA AAATTATTAA AATCAAAAAA CGTATAATAT CAGATATTCT	1380
AAAACCTTGA TACTGTACGT TTTATCATAG AAATTTTTAC TTTATTTTCT CATCAAATGA	1440
GATTGCGATC AATCTCTTGT CTTACTTGGC TTTCTTCTTC GCTTCTTCA TTTGTTAGC	1500
CATACGTTTC ATGGACTGTT TCATGGAAA TTCACCAATT TTACCTTCA AACGCCACC	1560
AAACATCTGG CTCATATCTG GCATTCCTGC TCCTCCGAGA GCTGATAAGT CAGGCATACC	1620
GCCTTGTCCC ATCATTCCTT CAAGGGCAGA CATATCCATT CCTCCCATAT TTGGCATATT	1680
TTTACCAAGG TTATTTGGAT TAATCCCCAT TTGCTTCATC ATTTTATTCA TATCCCCAGA	1740
CATAACACCC TGCATGAGCT GTTACGCCTG GTTAAAGTCC TTGATGAATT TATTGACTTC	1800
GACGAATGTA TTTCCAGAAC CAGCAGCAAT ACGACGGCGA CGGCTTGGAT TTAACAAATC	1860
TGGGTTTTCA CGCTCTTCAG GTGTCATCGA AGACACAATG GCACGTTTAC GAGCAATCTG	1920
GCGTTCATCC ACCTTCATGT TTTGAAGGGC TGGATTGTTG GCCATACCTG GAATCATCTT	1980
GAGCAAGTCT TCCATCGGCC CCATATTTG CACCTGATCT AATTGATCGA TGAAATCATT	2040
AAAATCAAAG GTGTTTCTGC GCATCTTCTC AGCCATTCA AGGGCTTTTT GTTCATCGTA	2100
TTCCCTGAGAA GCTTCTCAA TCAAAGTGAG CATATCCCCC ATACCAAGGA TACGGCTAGA	2160
CATGCCGTCT GGGTGGAGG TTTCATGTC CGTAATCTT TCACCTGTAC CAGTGAACCTT	2220
GATTGGTTTT CCAGTAATGT GACGAACAGA CAGAGCAGCA CCACCACGAG TATGCCATC	2280
AATCTGGTA AGGATGACCC CAGTCACTTC CAACTGAGCA TTAAACTCAC GCGAACATT	2340
GGCTGCTTCC TGACCAATCA TAGCATCAAC GACAAGCAAG ATTTCATTTG GTTGAGCCAA	2400
TGCTTCAACA TCACGAAGCT CATTCAATGAG GAGCTCATCA ATCTGCAAAC GACCCGCAGT	2460
ATCAATCAAG ACATAGTCGT TATGATTAGT TTGGGCTTGC TCCAAACCTT GACGTACAAT	2520

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CTCAACAGCT GGTACTTCTG TTCCAAGTGC AAAGACAGGC ACATCAATCT GTTGTCCCAA	2580
GGTCTTAAGC TGGTCAATGG CAGCTGGACG ATAAATATCC GCCGCAATCA TCAAAGGACG	2640
AGCATTTCCT TCTTTCTTGA GTTGTGCGC CAATTTACCA GCAAAGGTTG TTTTACCCAGC	2700
CCCTTGTAAA CCAACCATCA TGATGATGGT TGGAAATCTTA GGTGACTTGA TAATTTCTGC	2760
CGTATCAGAA CCTAAACCG CTGTCATTC CTCATCAACG ATTTAATAA TCTGTTGCC	2820
AGGATTAAGT GTATCAATGA CCTCATGCC GACTGCACGC TCACGAACCT TCTTGATAAAA	2880
GTCCTTACA ACAGGCAAGG CAACGTCGGC CTCGAGCAAG GCCAAGCGAA TTTCTTGGT	2940
TGCCTTGG ACATCAGATT CAGAGATTT TCCTTTTTA CGTAGATTT TAAAGACGTT	3000
CTGCAAACGT TCTGTTAACC TTTCAAATGC CATTTCCTT CCTCTTATTC TCTATTATCA	3060
ATGCTTGTAA AAATTCCTAT CTGCTCTGC AGAAAGTCAT CTTGGGATA GCGCTCCAAA	3120
ATCTGATCAA AAATCTGACT GCGGACAATA TAGTCCGAGT ACATGTGCAA TTTCATCTCA	3180
TAATCTCCA GAATCTTTTC TGTCGCTTG ATATTGTCA AGACAGCCTG ACGACTGACA	3240
CCGAACCTCCT CGGCAATTTC AGCAAGGCTG TAATCATCAG CGTAGTAGAG CTCGATATAA	3300
TTCAATTGCT TATCTGTCAA AAGCGCCGCA TAAAATTCAA AGAGCGCATT CATAKGATTG	3360
GTTCATTCGA TTTCATAAAC TTTTATTATA CCAAAATTA GCCTAATCTA CCACACTAGG	3420
AAGCCGATCC AAGAAGATAG ATAGCTAAAT TTGAAAAGA CATGAGCCTA GCCCCAAGTA	3480
ATTTCCAATT GATACTGGC AAAGGGATGT CCCTCTTGAT TTGTTAGTTG ATAATCTAGT	3540
TCAATCTTT GCCTATCAAC TTGATAATGG CTCGTTGGA TGATAAACTC CTGCATGCC	3600
ATAGGTGTAG GAATATAGGC TAAACTATCG CTATCCTTA GAAAGCGCAT AATGGTCTTG	3660
GGATTAGAAA ATCGGCTCAT CACAAGTTCT TGACCATGAA ATTAAATCAC TACTTTTCC	3720
TTTCCTCAT TATAGAAAAG CAGGTAGCTA TAATCTCTT TTTCATGCAC TTCCACATCA	3780
TAAAGCTGGT CAATCACTTC CAACTGCTCA TCAAACGTAA TGTTGTTTCG CATCCGAATC	3840
TTCACATCAG GCCCTTTTC TTGTCCTTG TCCTACTATT TTACCAAAAA GAGCAGGATT	3900
TTGCTATAAT GGTCATATGA ACGAAAAAGT ATTCCGTGAC CCTGTTACACA ACTACATCCA	3960
TGTCAATAAT CAAATCATCT ATGACTTGAT TAATACAAA GAATTCAGC GTTGTGCGCC	4020
GATCAAACAA CTGGAAACTT CCAGTTATAC CTTCCACGGT GGAGAACACA GTCGCTTCTC	4080
TCACTGTCTA GGAGTCTATG AAATTGCACG ACGCATCACA GAGATTTCG AAGAAAAATA	4140
TCCTGAGGAA TGGAATCCTG CCGAGTCTCT CTTGACCATG ACCGCTGCTC TCCTACACGA	4200
CCTTGGGCAT GGTGCCTACT CCCATACCTT TGAACATCTC TTTGATACAG ACCATGAAGC	4260

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CATTACTAG GAGATTATTC AAAATCCTGA GACAGAGATT CACCAAGTCC TGCTACAAGT	4320
GGCACCTGAT TTCCCAGAAA AGGTGGCCAG TGTCATTGAC CATACTATC CTAATAAGCA	4380
GGTCGTGCAG CTCATTTCTA GTCAGATTGA CGCAGATCGC ATGGACTATC TCTTGCGCGA	4440
CTCCTATTT ACAGGGAGCAT CCTATGGGG AATTGACCTG ACTCGAATCC TCCGAGTCAT	4500
TCGTCCTATC GAAAATGGTA TCGCCTTTCA GCGCAATGGC ATGCACGCCA TCGAAGACTA	4560
CGTCCTCAGT CGCTACCAGA TGTACATGCA GGTTTATTC CACCCCGCAA CACGCGCCAT	4620
GGAAGTTCTC CTACAGAACG TTCTCAAACG CGCCAAGGAA CTCTATCCTG AGGACAAGGA	4680
TTTCTTGCC CGAACCTCTC CACACCTCCT GCCTTTCTC GAAAAAAATG TGACCTTGAC	4740
TGACTATCTG GCTCTGGATG ATGGCGTGTATG GAATACCTAC TTCCAGCTTT GGATGACCAAG	4800
TCCTGACAAG ATTCTTGCGAG ATTATCGCA TCGCTTGTC AACCGCAAGG TCTTTAAATC	4860
CATTACCTTT TCACAAGAGG ACCAAGATCA ACTTACTAGC ATGAGAAAAT TGGTTGAGGA	4920
TATCGGCTTT GATCCCGACT ACTACACTGC CATTCTAAAG AACTTTGACC TCCCTTATGA	4980
TATCTATCGT CCCGAATCTG AAAACCCACG GACACAGATT GAGATTTAC AAAAAAAATGG	5040
AGAACTGGCC GAACTCTCTA GCCTGTCTCC TATCGTCCAA TCCCTTGCTG GCAGTCGCCA	5100
CGGAGATAAT CGCTTTTATT TTCCAAAAGA AATGTTGGAC CAAACAGCA TCTTGCAAG	5160
CATTACCCAG CAATTCTTAC ACTTGATTGA GAACGATCAT TTTACCCAA ATAAAAACTA	5220
GAAGAGGAAA TTATGAGTA TTAAACTAAT TGCCGTTGAT ATCGACGGAA CCCTTGTC	5280
CAGCCAAAAG GAAATCACTC CTGAAGTTTT TTCTGCCATC CAAGATGCCA AAGAAGCTGG	5340
TGTCAAAGTC GTGATTGCAA CTGGCCGCC TATCGCAGGC GTTGCACAAAC TTCTAGACGA	5400
CTTGCAGTTG AGAGACGGAGG GGGACTATGT GGTAACTTC AACGGTGCCC TTGTCCAAGA	5460
AACTGCTACA GGACATGAGA TTATCAGCGA ATCCTTGACT TATGAGGATT ATCTAGATAT	5520
GGAATTCCCTC AGTCGCAAGC TCGGTGTCCA CATGCATGCC ATTACCAAGG ACGGTATCTA	5580
TACTGCAAT CGCAATATCG GAAAATACAC TGTACACGAA TCAACCCCTCG TCAGCATGCC	5640
TATCTTCTAC CGTACCCCTG AAGAAATGGC TGGCAAAGAA ATTGTTAAAT GTATGTTAT	5700
CGATGAACCA GAAATTCTCG ATGCTGCGAT TGAAAAAAATT CCAGCAGAAAT TTTACGAGCG	5760
CTACTCCATC AACAAATCTG CTCCCTTCTA CCTCGAACTC CTTAAAAAGA ATGTAGACAA	5820
GGGTTCAAGCC ATTACTCACT TGCGCTGAAAA ACTCGGATTG ACCAAAGATG AAACCATGGC	5880
AATCGGTGAT GAAGAAAATG ACCCGTCCAT GCTGGAAGTC GTTGGAAACC CCGTTGTCAT	5940
GGAAAATGGA AATCCAGAGAA TCAAAAAAAAT CGCCAAATAC ATCACCAAA CAAATGACGA	6000
ATCCGGCGTT GCCCATGCCA TCCGAACATG GGTACTGTAA AAGTATCATT TTTCAATAAG	6060

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AATTGATTAG CAATAAAATC CAATGAATTT TTTTAGCAAA CTATTTAATT TAAAACAAAA	6120
TAATCATAAT AGAGACACAA ATTCTGATTG TAACAATTTT TACCTAACG AATTAGAATG	6180
TGGCCTTACT CCTGGGCAAC TCATACTCAT AGATTGGACT CAAAAAAACAG GGAGAAATTA	6240
TAATTTCCA AGATATTTTA AATACTCTCT TCAAATTGAC CCTGAATCTA CACACAATCA	6300
ATTATACAAA TTAGGATACT TCACTAAAAA TAAGACTTTA TCATATCTTA CAGTAGTAGA	6360
ATTAAAAACT ATATTATCTA AACATAATTT AGCTACTTCT CGAAAAAAAG CAGAATTAAT	6420
TACAAGAATA ATTAATAATG TTAACATTGA CAATTTAGAT ATTCCGTTCG AATTAAACT	6480
AACAAAAGAA GCACAAAATC TTATTATCGA ACATAGTGAC TATATCAAAG CATACTATGA	6540
TAAAGACATA ACTATGGAAG ATTATTGTAA AGAAAAAAAC AATATCTCTT TTAAAGCAAC	6600
TTTTGGTGAT ATAAAATGGA GTCTCTTAA TAAACAGCT CATAGGAATA CTGTATCAGG	6660
AGATTTGGA TGCTTATCTA ACACACGAA GGCTCAGGGAGACATTGG AACAGAAGG	6720
TAATATTTAA CATGCTTAA TATATTACAT AGAATCTTG ATAATTACTA TTTCAGGATT	6780
AGAAAACAAT TTTCAGCCA CTGATTATCC ACTATATTAT CCCGATTGCA TACCTGACTA	6840
CTCACTAAAA CATATTCAA CATTAATGGA ATCATTATCT GATGACGATT ATGATTTCGC	6900
TTTTGATGAA GCATTATTC GCTTCTCAAT TTTGAATGCA AATCATTTT TATCTAAGGA	6960
AGATATTGAC TATTTAAGAG TTAATTAC TCGTTCCACT GCTGAAGAAA TAAACAATTA	7020
CTTAAAGAAA TATGAATGTT ATAGTCCTTT AAATAATTAA GAACATTGACG ATTTGAAATA	7080
AATTGACTAT ACAAAACATT ATATACTCGA TATAGTCTCA ATTTTATCTG ATGATTGCC	7140
AAATTTTCA ATAATAAAAC GCATAATATT ATGGAGACAA TCCCCTATAT TATGCGTTCT	7200
TTTAATATCA AAGACTTTT GACAAACTTC TTTGATATCT AATTACATGC CCCCTGCAGG	7260
AATCGAACCT GCAACTACTC CTTAGGAGGG ACTTGTATA TCCATTGAAAC TAAGGGAGCT	7320
AGATAAAAAC TCTGCTAAAT GACCAGAGTT TTTTAGTCGA ATTAACGACG GATTCTTTG	7380
ATACGAGCTG CTTTACCTTG AAGAGCACGC AAGTAGTACA ATTTCGCAGC ACGTACTTTA	7440
CCGTAACGAA CAACTTCGAT TTTTCAACA CGTGGAGTGT GGATTGGAA GATACGCTCA	7500
ACACCTACAC CGTTAGAGAT TTTACGAACT GTGTAGTTT CTGAGATTCC AGCACCTTA	7560
CGTGCAGATAA CAACACG	7577

(2) INFORMATION FOR SEQ ID NO: 47:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 4945 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double

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(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 47:

CCTCGCTGAT GATTGGTGCT GTTTTATTTG CTGGTCCAGC CTTGGCTGAA GAAACTGCAG	60
TTCCTGAAAA TAGCGGA _n CT AATACAGAGC TTGTTTCAGG AGAGAGTGAG CATTGACCA	120
ATGAAGCTGA TAAGCAGAAT GAAGGGGAAC ATGCTAGAGA AAACAAGCTA GAAAAGGCAG	180
AAGGAGTAGC GATAGCATCT GAAACTGCTT CGCCAGCAAG CAATGAAGCT GCAAACACTG	240
AAACTGCAGA ACCAGCTAGC GCAGCTAAC CAGAGGAAAA AGCAAGTGAG GTGGTTGCAG	300
AAACACCATC TGCAGAAGCA AAACCTAAGT CTGACAAGGA AACAGAAGCA AAGCCCAGAAG	360
CAACTAACCA AGGGGATGAG TCTAAACCAAG CAGCAGAAGC TAATAAGACT GAAAAAGAAG	420
TCCAGGCCAGA TGTCCTAAA AATACAGAAA AAACATTAAA ACCAAAGGAA ATCAAATTAA	480
ATTCTTGGGA AGAATTGTTA AAATGGGAAC CAGGTGCTCG TGAAGATGAT GCTATTAACC	540
GCGGATCTGT TGTCCTCGCT TCACGTCGGA CAGGTCATT AGTCAATGAA AAAGCTAGCA	600
AGGAAGCAAA AGTTCAAGCC TTATCAAACA CCAATTCTAA AGCAAAAGAC CATGCTTCAG	660
TTGGTGGAGA AGAGTTCAAG GCCTATGCTT TTGACTATTG GCAATATCTA GATTCAATGG	720
TCTTCTGGGA AGGTCTCGTA CCAACTCCTG ACGTTATTGA TGCAGGTCAC CGTAACGGGG	780
TTCCCTGTATA CGGTACACTC TTCTTCAACT GGTCTAATAG TATTGAGAT CAAGAAAGAT	840
TTGCTGAAGC TTTGAAGCAA GACGCAGATG GTAGCTTCCC AATTGCCCGT AAATTGGTAG	900
ACATGCCAA GTATTATGGC TATGATGGCT ATTTCATCAA CCAAGAAACA ACTGGGAGATT	960
TGGTTAAACC TCTTGGAGAA AAGATGCGCC AGTTTATGCT CTATAGCAAG GAATATGCTG	1020
CTAAGGTAAA CCATCCAATC AAGTATTCTT GGTACGATGC CATGACCTAT AACTATGGAC	1080
GTTATCATCA AGATGGTTTG GGAGAATACA ACTACCAATT CATGCAACCA GAAGGAGATA	1140
AGGTTCCGGC AGATAACTTC TTGCTAACT TTAACTGGGA TAAGGCTAAA AATGATTACA	1200
CTATTGCAAC TGCCAACCTGG ATTGGTCGTA ATCCTTATGA TGTATTGCA GGTTTGGAAAT	1260
TGCAACAGGG TGGTCCCTAC AAGACAAAGG TTAAGTGGAA TGACATTAA GACGAAAATG	1320
GGAAATTGCG CCTTTCTCTT GGTTTATTTG CCCCCAGATAC CATTACAAGT TTAGGAAAAA	1380
CTGGTGAAGA TTATCATAAA AATGAAGATA TCTTCTTTAC AGGTTATCAA GGAGACCTA	1440
CTGGCCAAAA ACCAGGTGAC AAAGATTGGT ATGGTATTGC TAACCTAGTT GCGGACCGTA	1500
CGCCAGCGGT AGGTAATACT TTTACTACTT CTTTTAATAC AGGTCAATGGT AAAAAATGGT	1560
TCGTAGATGG TAAGGTTTCT AAGGATTCTG AGTGGAAATTA TCGTTCAGTA TCAGGTGTT	1620

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TTCCAACATG GCGCTGGTGG CAGACTCAA CAGGGAAAA ACTTCGTGCA GAATATGATT	1680
TTACAGATGC CTATAATGGC GGAAATTCCC TAAATTCTC TGTTGATGTA GCCGGTAAGA	1740
CAGATCAGGA TGTGAGACTT TATTCTACTA AGTTAGAAGT AACTGAGAAG ACCAAACTTC	1800
GTGTTGCCA CAAGGGAGGA AAAGGTTCTA AAGTTTATAT GGCATTCTCT ACAACTCCAG	1860
ACTACAAAATT CGATGATGCA GATGCATGGA AAGAGCTAAC CCTTTCTGAC AACTGGACAA	1920
ATGAAGAATT TGATCTTAGC TCACTAGCGG GTAAAACCCT CTATGAGTC AAACTATTTT	1980
TCGAGCATGA AGGTGCTGTA AAAGATTATC AGTTTAACCT AGGACAATTAA ACTATCTCGG	2040
ACAATCACCA AGAGCCACAA TCGCCGACAA GCTTTCTGT AGTAAACAA TCTCTAAAA	2100
ATGCCCAAGA AGCGGAAGCA GTTGTGCAAT TAAAGGCAA CAAGGATGCA GATTCTATG	2160
AAGTTATGA AAAAGATGGA GACAGCTGGA ATTACTAAC TGGCTCATCT TCTACAACTA	2220
TTTATCTACC AAAAGTTAGC CGCTCAGCAA GTGCTCAGGG TACAACCAA GAACTGAAGG	2280
TTGTTAGCAGT CGGTAAAAT GGAGTTCGTT CAGAAGCTGC AACACAAACC TTGATTGGG	2340
GTATGACTGT AAAAGATACC AGCCTACCAA ACCACTAGC TGAAAATATC GTTCCAGGTG	2400
CAACAGTTAT TGATAGTACT TTCCCTAAGA CTGAAGGTGG AGAAGGTATT GAAGGTATGT	2460
TGAACGGTAC CATTACTAGC TTGTCAGATA AATGGTCTTC AGTCAGTTG AGTGGTAGTG	2520
TGGATATTG TTTGACCAAG CCACGTACCG TTGTTAGATG GGTCTGGAT CATGCAGGAG	2580
CTGGTGGTGA GTCTGTTAAC GATGGCTTGA TGAACACTAA AGACTTGAC CTTTATTATA	2640
AAGATGCAGA TGGTAGTGG AAGCTAGCTA AGGAAGTCCG TGGTAACAAA GCACACGTGA	2700
CAGATATCAC TCTTGATAAA CCAATCACTG CTCAAGACTG GCGCTTGAAT GTTGTCACTT	2760
CTGACAATGG AACTCCATGG AAGGCTATTC GTATCTATAA CTGGAAAATG TATGAAAAGC	2820
TTGATACTGA GAGTGTCAAT ATTCCGATGG CCAAGGCTGC AGCCCGTTCT CTAGGCAATA	2880
ACAAGGTACA AGTTGGTTT GCAGATGTAC CGGCTGGAGC AACTATTACC GTTTATGATA	2940
ATCCAAATTC TCAAACCTCG CTCGCAACCT TGAAGAGCGA AGTTGGAGGA GACCTAGCAA	3000
GTGCAACCATT GGATTGACA AATCAATCTG GTCTTCTTTA TTATCGTACC CAGTTGCCAG	3060
GCAAGGAAAT TAGTAATGTC CTAGCAGTTT CCGTCCAAA AGATGACAGA AGAATCAAGT	3120
CAGTCAGCCT AGAAACAGGA CCTAAGAAAA CAAGCTACGC CGAAGGGGAG GATTTGGACC	3180
TTAGAGGTGG TGTTCTCGA GTTCAGTATG AAGGAGGAAC TGAGGACGAA CTCATTGCC	3240
TAACTCACGC AGGTGTATCA GTATCAGGTT TTGATACGCA TCATAAGGGA GAACAGAAC	3300
TTACTCTCCA ATATTTGGGA CAACCGTAA ATGCTAATTT GTCAGTGACT GTCACTGGCC	3360

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AAGACGAAGC AAGTCGAAA ACTATTTGG GAATTGAACT AAGTCAGGAA CCGAAAAAAG	3420
ATTACCTAGT TGGTGATAGC TTGACTTGT CTGAAGGACG CTGGCAGTG GCTTATAGCA	3480
ATGACACCACAT GGAAGAACAT TCCTTTACTG ATGAGGGAGT TGAAATTTCT GGTTACGATG	3540
CTCAAAAGAC TGGTGTCAA ACCTTGACGC TTCATTACCA AGGCCATGAA GTTAGCTTG	3600
ATGTTTGGT ATCTCCAAA GCAGCATTGA ACGATGAGTA CCTCAAACAA AAATTAGCAG	3660
AAGTTGAAAGC TGCTAAGAAC AAGGTGGTCT ATAACTTGTC TTCATCAGAA GTAAAAGAAC	3720
CCTTCTGAA ACCAATTGAA GCGGCCGAAC AAGTGTGAA AGACCATGAA ACTAGCACCC	3780
AAGATCAAGT CAATGACCGA CTAAATAAT TGACAGAACG TCATAAAGCT CTGAATGGTC	3840
AAGAGAAATT TACGGAAGAA AAGACAGAGC TTGATCGCTT AACAGGTGAG GTTCAAGAAC	3900
TCTTGGCTGC CAAACCAAAC CATCCTTCAG GTTCTGCCCT AGCTCCGCTT CTTGAGAAAA	3960
ACAAGGCCCT GTTGAAAAA GTAGATTTGA GTCCAGAAGA GCTTACAACA GCGAAACAGA	4020
GTCTAAAAGA TCTGGTTGCT TTATTGAAAG AAGACAAGCC AGCAGTCTTT TCTGATAGTA	4080
AAACAGGTGT TGAAGTACAC TTCTCAAATA AAGAGAAGAC TGTCATCAAG GGTTGAAAG	4140
TAGAGCGTGT TCAAGCAAGT GCTGAAGAGA AGAAATACTT TGCTGGAGAA GATGCTCATG	4200
TCTTTGAAAT AGAAGGTTTG GATGAAAAAG GTCAAGATGT TGATCTCTCT TATGCTTCTA	4260
TTGTGAAAT CCCAATTGAA AAAGATAAGA AAGTTAAGAA AGTATTTTC TTACCTGAAG	4320
GCAAAGAGGC AGTAGAATTG GCTTTGAAAC AAACGGATAG TCATGTTATC TTTACACCAAC	4380
CTCACTTTAC TCATTATGCC TTTGTTATG AATCTGCTGA AAAACCAACAA CCTGCTAAAC	4440
CAGCACCCACA AAACACAGTC CTTCACAAAC CTACTTATCA ACCGACTTCT GATCAACAAA	4500
AGGCTCCTAA ATTGGAAGTT CAAGAGGAAA AGGTTGCCCT TCATCGTCAA GAGCATGAAA	4560
ATACTGAGAT GCTAGTTGGG GAACAACGAG TCATCATACA GGGACGAGAT GGACTGTTAA	4620
GACATGTCTT TGAAGTTGAT GAAAACGGTC AGCGTCGTCT TCGTTCAACA GAAGTCATCC	4680
AAGAAGCGAT TCCAGAAATT GTTGAAATTG GAACAAAAGT AAAAACAGTA CCAGCAGTAG	4740
TAGCTACACA GGAAAAACCA GCTCAAATA CAGCAGTTAA ATCAGAAGAA GCAAGCAAC	4800
AATTGCCAAA TACAGGAACA GCTGATGCTA ATGAAGCCCT AATAGCAGGC TTAGCCAGCC	4860
TTGGCTTGC TAGTTTAGCC TTGACCTTGA GACGGAAAAG AGAAGATAAA GATTAATAT	4920
CGAAAAATCT TGTGAAATCT TTCCG	4945

(2) INFORMATION FOR SEQ ID NO: 48:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 25002 base pairs
 - (B) TYPE: nucleic acid

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(C) STRANDEDNESS: double
 (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 48:

GACAACTCAA GTAGCTTTT CTTATTTGA AAAAGGAGAT CAGAGTTAA CTATGTCAGA	60
AAAATCACAA TGGGGTCTGA AACTTGGTTT TATTCTAGCA TCTGCTGGCT GGCCATCGGG	120
CTTGGTCCG TTTGGAAGTT TCCCTACATG ACTGCTGCTA ATGGCGGTGG AGGCTTTTA	180
CTAACATTTTC TCATTTCCAC TATTTTAATC GGTTTCCCTC TCCTGCTGGC TGAGTTGCC	240
CTTGGCCGTA GTGCTGGCGT TTCCGCTATC AAAACCTTG GAAAAGTGGG CAAGAATAAC	300
AAGTACAAC TTAGTGGTTG GATTGGCGCC TTTGCCCTCT TTATCCTCTT ATCTTTTAC	360
AGTGTATCG GAGGATGGAT TCTAGTCTAT CTAGGTATTG AGTTGGGAA ATTGTTCCA	420
CTTGGTGGAA CGGGTGATTA TGCTCAGTTA TTTACTTCAA TCATTTCAA TCCAGCCATT	480
GCCCTAGGAG CTCAAGCGGC CTTTATCCTA TTGAATATCT TCATTGTATC ACGTGGGTTT	540
CAAAAAGGGA TTGAAAGAGC TTCGAAAGTC ATGATGCCCT TGCTCTTAT CGTCTTTGTT	600
TTTATCATCG GTCGCTCTCT CAGTTGCCA AATGCCATGG AAGGGGTTCT TTACTTCCTC	660
AAACCAGACT TTTCAAAACT GACTAGCACT GGTCTCCTCT ATGCTCTGGG ACAATCTTC	720
TTTGCCTCT CACTAGGGGT TACAGTCATG TTGACCTATG CTTCTTACTT AGACAAGAAA	780
ACCAATCTAG TCCAGTCAGG AATCTCCATC GTAGCCATGA ATATCTCGAT ATCCATCATG	840
GCAGGCTCTAG CCATTTCCA AGCTCGATCC CCCTTCATAA TCCAGTCTGA AGGGGGACCC	900
AGCCTGCTCT TTATCGTCTT GCCTCAACTC TTTGACAAGA TGCTTTGG AACCATTTTC	960
TACGTCCTCT TCCTCTTGCT CTTCTTTT GCGACAGTC CTTTTCTGT CGTGATGCTG	1020
GAAATCAATG TAGACAATAT CACCAACCAG GATAACAGCA AACGTGCCAA ATGGAGTGT	1080
ATTTTAGGAA TTTTGACCTT TGCTTTGGC ATTCCTTCAG CCCTATCTTA CGGTGTCATG	1140
GCGGATGTTACATTTGG TAAGACCTTC TTTGACGCTA TGGACTTCTT GGTTCCAAT	1200
CTCCTCATGC CATTGGAGC TCTCTACCTT TCACTTTTA CAGGCTATAT CTTTAAAAG	1260
GCTCTGCAA TGGAGGAACCT CCATCTCGAT GAAAGAGCAT GGAAACAAGG ACTGTTCAA	1320
GTCTGGCTCT TCCTCTTCG TTTCTTCGTT TCGTCATTC AATCATCATC ATTGTGGTCT	1380
TCATTGCCA ATTTATGTA TCAAAAGGA CTTGAGTAGT GAACTCAGGC CCTTCTTTT	1440
TATGGATGGC TAACAATCAA TTCCAAACCT TGCCCTTCCA GAGTCCAAGC TTCAACATCA	1500
CTTGGTAGGA TAAAGTGGCT GCCTTTTGA ATTGGATAAT TTTTCCCCGTC AACAGTTAGC	1560

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TGACCTTGAC CAGCCAAGAC ACTCAATAAG CTGTAGTCAG CTGTCTTTTC AAAGTCAACT	1620
TTTCCAGTAA TTTCCCACCT GTAAACTCGG AAGAAATCAT TAGATACAAG GAGAGTGGAA	1680
CGCAAATCAT CTGCTTTAAC AGTTACAGGA CGGCTATTG CTGGCTCACC AATGTTCAAG	1740
ACATCGATGG ATTTTCAGA ATGAAGTTCAGA CGCAAGTTGC CTTTGTCAATC CTTGCGGTCA	1800
AAGTCATAGA CGCGATAGGT GGTATCGCTA GACTGCTGGG TTTCAAGGAT TAAGATAACC	1860
CCCCCGATAG CGTGCATAGT CCCGCTTGGT ACATAGAAGA AATCTCCAGC CTTAACAGGG	1920
ACTTTGGTCA ACAAGTCATC CCAGTTCTTG TCCTCGATTT GCTGGCGGAG TTCTTCTPTT	1980
GACTTGGCAT TGTGACCGTA GATAATCTCT GAACCTTCAT CCGCTGCGAT AATGTACCAAG	2040
CATTCTGTTT TTCCGAGTTC GCCTTCATGC TCGAGTCCAT AAGCATCGTC TGGGTGAAC	2100
TGGACACTGA GCCAGTCGTT GGCACTCGAGG ATCTTGGTCA AAAGTGGAAA TACAGGTTCT	2160
GGACGATTGC CAAATAATTCC ACCGTGTTCC GCATACAAAG TAGCAAGATC TGTTCCCTCG	2220
TAACGACCAT TGGCAACTTT AGAGACTCCA TTTGGATGGG CTGAGATGCC CCAATATTCT	2280
CCGATTTTTT CACTTGGGAT GTCGTAGCCA AACTCATCAC GTAGCTTGGC TCCACCCAG	2340
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GCTCCGTTTC CTACATTATC TCGATCTACA TAGATAAAGC CATAGCGCTT ATTCAATTCC	2640
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TAGTCATCTG CTACATAACC ATTCTCATCC GGTGTATCCA TAGCACCGAG TCCATTTCT	2820
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GAAGAAATCC AAACCATCAA AACACTTTTA AAAGACTCTC GTACAGCTAA ATATCATAAA	2940
CGCCTCAAA TCGTTCTATA GTAAAATGAA ATAAGAACAG TACAAATCGA TCAGGACAGT	3000
CAAATCGATT TCTAACAAATG TTITAGAAGT AGGGGTGTAC TATTCTAGTT TCAATCTACT	3060
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CAAGGTGTAC TGTATAGAT TCAATACACT ATAGACTGTA ATCAAACAAAC GATTTGGCGA	3240
AATGTAAAAA AATATGAGGA GTTCGGACTC GACTCTCTCC TTCAAGAAC ACGTGGTGGT	3300
CGTAACCATG CATATATGAC AGTTGAGGAA GAGAAAGCCT TTCTTGCCTT CCATTTGAAG	3360

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GCTACAGAGG CAGGAGAATT TGTACAATT GATGCCATT TTCAGGCTTA TAAAAAGGAG	3420
TTAGGTCGTT CCTACACACG TGATGCCCTC TATCAACTGT TGAAGCGCCA TGGTTGGCGA	3480
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AAAAATAAAA TCTCAATCCA AGAAGGCAAG AAAGCGTTT AAAATAGTA GACGTTTCG	3600
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GCAATCCCCA CATCTGCTAC ACTAAGGGCA GGAGCGTCAT TGATACCGTC CCCAACAAAG	7020
GCTACTTTCG CTGACTGTTG CAGTTTATGG ATTTCATGGG CTTTTCTTC TGGCAAGACG	7080
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TTTTCTAACG	TTGCTTCAC AATTTCTGTC ATAGTCTCCA CCTACTCTAC AATCATCTG	8940
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TCCACTACAT	ACTCTCCCC CATTGGCAGG TTGCGATGTA CACCAAAATC TGGAAAACA	9060
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AGTCCGATAG	CTATTCTTC	ACTGTGCATG	TTCGTCATAT	CCGTGAGCAG	ATAGAGCTCT	13860	
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GAATCTCCAA TATCTCGATT CTGCTCTTGC GGTCAATGGA AGCGAAGGGC TCATCCAAGA	15300
GATAGACCCCT AGGATTCTAG GCAAAGAGGA CAGCCAGCGC TGCTTTTGC TTTTCCCCAC	15360
CTGATAAGTG ATGGATGAGA CGGTGCAAGA TGTCCCTGCA ACGACATTGC TGGACAAACCT	15420
CTGCTATTT AGAATCAATT TCCTGAAGGT GATAGCCGAT ATTTTCCATG GTAAAAACCA	15480
ACTCCTCAAA CAAGCTCTCC ATGGTAAATT GATGATTAGG ATTTTGCAAG AGAATACCAA	15540
CCGTCTGGAC ACGTTGACG ATAGAAAGCT GACTGACCTC GCTCCCATCT ATCAGGACTT	15600
GACCGCTATA GGGAGAGAA CTAACCTGGG CAATCATTG AAAGAGGCTG GATTTCCAG	15660
ACCCACTACT CCCACTAAC AAGGTAAGG CTTGCGCATG AAAAGTAAA TCAACGGCT	15720

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CAGAGAAAGAT	TGGGGACTGA	ATCGCTCGTA	GTTCCAGACC	CATCTATGCT	TTTCCTCCAG	15780
TTGCAAACGT	ATGATAGAGT	TTGACAATGG	CACGAACCAA	GATGGTACAG	AAGAAATAAA	15840
CAGAAATAAA	ACGTACCAACA	ACCAAGGAAA	GGACAAACGG	AAGGGAAAAG	GGTAGTAAC	15900
CTAACCTTAAT	GTATTCTAG	ACAAAGCTAA	CAAGCGTAAT	CCCAACTA	TTAGCGTTA	15960
GAGAGAGCCA	ACTTCATAG	CGATTCTTAG	TTACGATAAA	ACCAAATTCA	CTTCCCAAAC	16020
CTTGAACAAA	GCCAGACAAA	AGAGCTCCTA	GACCAAATTG	GCTACCATAA	AGGACTTCAG	16080
CAAGCCGCAGC	TAGCACTTCT	CCAATCGTTG	CACTTCCGAC	TCTCGGAACA	AAGATGGCAG	16140
CAATGGGCAGC	AGCCATACAC	CAGAGACCGA	AGAGGATTTC	ATTGGCAAAG	GCCTGCAAAC	16200
CAAGAGGTGT	TAAGAGTAGA	CTGAGAATAT	TATACACATA	TCCTGAACCA	ACGAAACCC	16260
CACCAAAAAA	GATAGACAAG	AAAGCAAGCA	AGATAACATC	TTTTAACTGC	CATTTTTCA	16320
ACATAAAAAA	CTCCTTTTTT	TAAGAGAAAAG	TGAGGCACTC	AAGAAGACCG	ACCTAAATAC	16380
TTTGTATAGC	AGACTGAATT	TAGAACAGTA	CACAAGAACAA	CTAAAATATT	TCTAGAAATT	16440
AATTTGAATT	TTCTAATTGA	TTTGTTCGCA	TCTTATTTC	ATCTACTATA	TCATCTTCAT	16500
CCAGTTCGT	AAAAGAAAAA	ACTCTAATTA	CAGATACAAA	TTAGAGTTCA	GCTTACAAGA	16560
TTAGACAGTT	CTTTTCGACA	TACGAAAAAA	ACATTTCACAA	TTTCCCTTCG	CCAGTCTTAA	16620
CTGTATCAGG	TTCAATGGGT	ATCATCTCAG	CCTAAAGCAC	CCCCAATGTC	TTTATTATTT	16680
AATTATGTGA	TTATTATAAC	ACACATTTA	TACTAGTTCA	AGAAATTGAA	CTGGAAATAC	16740
AGCCTTGAC	TCACAAAGAC	AGCAGATCTT	TCTTTTGCAA	AAAACAAATG	ACCTGTTGA	16800
TGAATTAGCC	ATTCAAGCTG	AATCTGGACA	TAGCTTTTTA	AAAAAGGAAA	ATCCTACTTA	16860
CTTAGAATCC	AAGGATAGAT	ATCTATTGTT	CACTCATTTTC	CCGAACAGTT	TTTTCTATAT	16920
TTTTTGATA	CGATATTGCC	GAAATGATTG	AAACGCCATC	CATATTGGTC	TTTATAATGT	16980
CTTTAATATG	TTTCGTCTGT	ATCCCACCAA	TTGCAACTAA	AGGCATTTGT	GGCAATAGTT	17040
TTCTCATCAA	TTCAAGACCT	TCATAACCTA	TAGTACCAACC	AGCATCATCC	TTTGACTGGG	17100
TACCAAATAC	AGGCCAACAA	CCTACATAAT	CTACATATTC	AACTTTGAT	TGTTGAAATT	17160
CTTCTCGTT	TCTTATAGAA	AGACCAATTA	TTTTATCTGG	CATCAATTTC	CTAATTCAT	17220
CAACACCAAT	ATCATCTTGA	CCTACATGTA	CGCCATCGGC	GTCAATTTC	ATTGCTAAAT	17280
CTATATCGTC	ATTAACGATA	AATGGAACAT	TGTATTTTTT	ACAAAGTTCT	TTAATTGGA	17340
TAGCTAGCTC	AAGTTTTCT	AAGCCTCTA	AAAGCACCCTC	ACCTTTTCT	CGAAATTGAA	17400
ATAAGGTTAT	ACCACCTTT	AAGGCTTCCT	CAACGACTGT	ATATAGATT	TTTCCTTGGC	17460
AAAGTAGTCGT	TCCACAAATA	AAATATAGTT	TTAGTAATT	TTATGAAAC	ATCTTACTTC	17520

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ACTCTTTGA ATTCCCTTAC ATCTTCATCT GTAATCTCGT ATAAGGCATT TATAAATTCA	17580
ACTTTAAATG TCCCAGGAAG ATGTCCATT GGACGTTTT CTGCTATTTC TCCAGCGATA	17640
TTGTAAACCA ACACTGCTGT TTTAATGAT TTCAATTCTT GACCTTTTC TAGTCCGATA	17700
AAGCTTGCTA CTACAGCTCC TAATAAGCAT CCTGTCCCAA TGACTTTCGG CATCATAGCA	17760
CTACCATTAT GAATCATTAC CACTTCTCCA TTAACAGCAA TGGCATCCAC TTCACCTGTT	17820
ACTACTATTG GAATATTGAA CTTCTCATTT GCTGCTAGAG CAATTTCGTC AATATTATCT	17880
ACGCCGCAC TATCTACTCC TTTAGATGCC ACATCTATTG CTACTAAAGA GGCAATCTCG	17940
CCAGCATTTTC CTCTAATCGC TGCTAGTTA TAATTGTTGA TTAGATCATC TGCTACTTTT	18000
TTTCTATATT CTCCGTCTC ACAGGCTACA GGATCTAAAA CTGCTGGGAC ATTATATTTC	18060
TCTGCAATTTC TCAGAGCAGC TTGGTATAAT TTCCAATTTCATCTGTCAA TGTCCTATG	18120
TTTATTAATA AACCCACCAGC ATACTTAAC AAATCCTCTA AATCTGCTGG AAACTCACTC	18180
ATGGCTGGTG AGGCCGCAG TGCTACTAAT CCATTTGCTG TGAAATTTC TACTACATCA	18240
TTGGTTATAC AAATGACCAA TGGTGTTTT TCTTTTAATA ATTTTAAACT TGTCAATTG	18300
AAATCCTTCC TTTTCACTTT ATACGATCTA CTAATTTCGA TTTATCTTA GTTGAGAATT	18360
TTTTCACTTT ACATTGAATG ATTTATACTC AATGAAAATC AAAGAGCAAA CTAGGAGGCT	18420
AACCGCAGGT TGCTAAAC ACTGTTTGAA GGTTGTTGGAT AGAACTGACG TGTTTGAAG	18480
AGATTTTCGA AGAGTCTTAC CTCATCAAAT TTGTAAATAT CATGAGCCTT CTCTAGACAT	18540
CGTAACCAAT ATCAAAAAAA GCTAATTCTA AAGCGACTGC TTGATTCCAG CGTTGCTGAA	18600
GTTCTGCAA ATCTTCTCGA TTTTTACCGA CACGATTGAG TTGCTCAACC AGAAATTGAA	18660
CCCACCTCTGC AAAGAAAGGA CCTCTGTGGA GATTGATCCA TTCCGAATGA ATATAGACTT	18720
CAGGTTAAAGC CAAATCTTA GAACCCCGAT CTAATAGAG ACCTTCTGCA ATGACCAGCA	18780
TGACCAAAAG ATGGCATAG TCTGATGAAG CCACCGCCGA ATACATTAGA TCCTGAAAGG	18840
CTTTTGTAC AGGGTGCAA GTCACTTCTA GATAGTCATT CTCTGCTACT TTTAACTCTT	18900
TAAAAGCCTT TTGGAAATAA CCATCTTCAT CTGCTTCAG AAAGCCTAGT TGCTTGCGAA	18960
AACGAACCTT GGATTCAAGT TTATCTGCGT GACTACGGAG GCACCCAGCA TGGATAAGAA	19020
GGCATCAAAG AAGTGATAAT CTTGAATCAG ATAGTCCTTT AAGACCTTAT TCTCAATTGT	19080
CCCCGCAAAA AGTCCCTAA CAAAACGATG ATTGATTGCA GCCTGCCAAT CCTCTGACT	19140
GCTTTTAAT AATTCTCCAA CAGTCAAACC TGGCTGAAAT GCATAGTCCT GTGTTCCAT	19200
ATTTACTTCT CCTCTCTTAA CTTGTTAGTA ATTAATAAAA CACCAAGAAA TATCAAGCAA	19260

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AATCGTAATT CCACCTGATC CTTTTAAAGC ACATCGAGAG CATTGCAGA GAGCTAACTA	19320
AACAAGCCTA TCCAGTTAT ATAAACAAAA AACTCCAATT ACAATCAAGA ATTAGAGTTG	19380
ACTTACAAGA TTAGACCGTT CATTCAACCA TACGAAAAAA CTGTTCACAT TTCCCTTCGC	19440
CAGTCTAAC TGTATCAGGT TCAATGGGT TTATCTCAGC CTAAGCACC CCAAATGTCT	19500
CTATTATTTA ACTACTGAAC CAGTATAGCA AAAATGAAA GCCCTAGCAA GATATTTGAC	19560
CGAAAAATAT CTTTATATAT AATATATTGA AACTAGATA GTACACCTCT ACTTATAAAA	19620
CATTGTTAGA AATCGATTTG ACTGTCCTGA TTGATTTGTC CTATTCTTAT TTCATTTTAC	19680
TATAGTTTC GATAGCAATT TATTCTTCA ATACACGAAG AAAACCTCC ACATTCACTG	19740
GAGGCAATCT GTTTATCAA TACAATTTA AGTCACGAGG GTCAACTGGG AAGGTTGGGT	19800
TGTATGGATT GTGACGGAGC TTGAAGTGTG TGACATCTC AATGGTCTGA GTTCCAGACA	19860
ATTGCATAAC TGTCTTCAAT TCCGCATTCA ACTGTTCAA GACTTGACCC ACACCGACAC	19920
TACCAACCGAG AGCCAAGCCA TAGATGACAG GGCGTCCAAT ACCAACCAAG TCTGCTCCTG	19980
ATGCCAAGGC TTTAAAGACG TGTGTGACCAC GACGAACACC AGAGTCAAAG ACAATCGGCA	20040
CACGTCTATC AACTGCTTCTT GCCACTTCTT GAAGCGACTC AAAGGCAGCT GGTCCACCGT	20100
CGATTTGACG ACCACCGTGG TTGGTTACCC AGATACCAGA AGCTCCTGCA GCAAGCGAAC	20160
GTTCAACGTC CTCACGGCAT TGTGGTCCCT TGACATACAC AGGAAGACCA GAGTATTCA	20220
CGATAAAATTC TACATCGCGT GGAGACAAGC GTTGTGTTAGC TGATTTGTAA ACMAACTCCA	20280
TTGATTTACC AGCACCTCTT GGCAGGTATT CTTCAACAAAT CGGCATGCCA ACTGGGAAGA	20340
CAAAACCCATT ACGCTTATCC ACTTCACGAT TCCCCCTAC AGTAGCATCT GCGTCAAGA	20400
CAATCGCTT ATAACCTTCA GCCTTCACAC GGTCCATGAT GTGGCGGTTG ATACCGTCAT	20460
CCTTACTAAA GTAAAATTGA AACCAATGAG GTGTCCCTG GAGGGCTTCA GAAATCTCTG	20520
GAAGGTCAAC AGTAGAGTAA GAACCTGGTG TATAAAGAGA ACCAAACTCA TGCACACCC	20580
GCGCAGTCGC CACTCCCCC TGTTCATTG CCAATTTATG AGCCGCAACA GGTGCCATAA	20640
TGATTGGAGA AGATAGTTT TCACCTGCAA ATTCAATCTC TGTACTTGGA TTTTCTACAT	20700
TGCAAAGTGT ATGAGGAACG ATGAGCTTGT GTTAAAGGC ACGGATATTC TCTCTTAAAG	20760
TGAAAGTATC TTCCGCCCA CTAGCGATAT AGCCAAATGC TGCTTTAGGA ATAACCTGTT	20820
GCGCCATGG CTCCAAATCA TAGGTATTGA TGAATCTAC ATGACCTTCT GCATTGCTTG	20880
TTTTGTATGA CATAAAATGT CCTCTTAAT AAGTAAGCGT TTACTTTGTC TATTACAAA	20940
ATATCTAAC TCTTTTCAA AACTTTAAA ATATTTGTT TGGAAATTTC AGAAATTAA	21000
TGTCTATGAT AAAATCCTT ATAACGGCAA TAAAAAATAG ATATTATCCA AAGAAGATTT	21060

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TAAGTGCTAC AATAACTGTA TTATTTCTAG ATGGGAGGTT CTATTTTGG ATTGATCCAT	21120
TGTTGAACAA TATCTACCAC TATATCAAAA GGCAATTCTTT CTGACCTTGC ATATTGCAGT	21180
TTGGGGAATT TTGGGATCCT TTCTGCTCGG TTAAATCGTT AGTATCATCC GACATTATCG	21240
AATCCTTGTG TTGGCGCAAG TAGGCACAGC CTACATTGAA TTGTCACGTA ATACGCCCT	21300
TTTGATTCAA CTCTTCTTC TCTACTTCGG TCTTCCCCGA ATCGGGATTG TCCTATCTTC	21360
AGAAGTCTGT GCAACGCTTG GGCTTGTCTT TTTAGGAGGC TCCTATATGG CAGAACCTTT	21420
CCGAAGTGGG CTGGAAGCCA TCAGTCAAAC CCAGCAGGAG ATTGGCCTCG CTATTGGTCT	21480
GACACCTCTA CAGGCTTTT ACTATGTGGT TCTTCCGAA GCAACAGCGG TGGCACTCCC	21540
CTCCTTTAGT GCCAATGTCA TTTTCTTAT CAAGGAAACC TCTGTTTCT CAGCAGTGGC	21600
TTTGGCCGAC CTCATGTACG TCGCCAAGGA TTTGATTGGT CTCTACTATG AGACAGACAT	21660
TGCGCTAGCT ATGTTGGTAG TTGCTTATCT AATCATGCTG CTACCCATCT CACTGGTCTT	21720
TAGCTGGATA GAAAGGAGGC TCCCCATGC AGGATTGGG AATCCAAGTA CTCTTCAAG	21780
GAAAATATCT CCTGAGAATC TTACAGGGAT TGCGCTTAC GATTGGATA TCCATCCTGT	21840
CTGTCCTCTT ATCCATGATG TTCAGAACAG TCATGGAAT CATCATGACC TCCCATTCTA	21900
GAATCATACG ATTTTAACA CGATTGTATC TGGAATTAT CCGTATCATG CCCCAGCTGG	21960
TGCTACTCTT CATCGTTTAC TTTGGCTTGG CTCGAAACTT TAATATCAAT ATCTCAGGTG	22020
AGACTTCAGC TATTATCGTT TTTACCCCTCT GGGGAACAGC TGAAATGGGA GACTTGGTAC	22080
GTGGAGCTAT CACTTCTCTC CCTAAACATC AGTTGAAAG TGGACAGGCA CTCGGCTTGA	22140
CTAATGTTCA ACTTTACTAC CACATCATCA TCCCACAAGT CTTAAGAAGA CTGCTACCGC	22200
AGGCTATCAA TCTTGTCACT CGGATGATTA AAACCACTTC ATTAGTTGTT TTGATTGGGG	22260
TTGTGGAAGT GACCAAAGTT GGACAACAAA TCATCGATAG CAATCGCCTG ACCATCCCAA	22320
CTGCTTCATT TTGGATTAT GGAACCATTC TAATCTTATA TTTCGCAGTT TGCTACCCCTA	22380
TTTCCAAACT ATCCACTCAC TTAGAAAAAC ATTGGAGAAA CTAAATGTCT GAAACTATCT	22440
TAGAAATCAA GGAACTAAAA AAATCCTCG GAGACAATCC CATCCTCCAA GGACTTTCTC	22500
TAGAAATCAA AAAAGGGAA GTTGTGTCA TCCTAGGGCC ATCTGGTTGT GGGAAAAGTA	22560
CCCTCCTTCG TTGCCTCAAC GGCTTAGAAA GTATTCAAGG TGGAGATATT CTTCTGGATG	22620
GTCAGTCTAT CGTTGAAAAT AAAAAGATT TTCACCTAGT TCGCCAAAAG ATTGGCATGG	22680
TCTTTCAAAG TTATGAACTC TTCCCCATC TGGATGTCTT ACAAAACCTC ATCCTAGGCC	22740
CTATCAAAGC TCAAGGAAGG GACAAGAAAG AAGTAACGGA AGAAGCTTTG CAATTACTAG	22800

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AGCGTGTGCG	TTTGCTGGAT	AAACAAACATA	GCTTTGCCG	TCAATTATCT	GGTGGACAGA	22860
AGCAACGTGT	TGCAATTGTC	CGTGCCTCC	TAATGCATCC	AGAAATCATC	CTTTTGACG	22920
AGGTGACTGC	TTCGCGTGGAT	CCAGAAAATGG	TGCGTGAGGT	GCTGGAACCTT	ATCAATGATT	22980
TGGCCCAAGA	AGGCCGTACC	ATGATTTAG	TAACCCACGA	AATGCGAGTT	GCCCAAGCCA	23040
TTACTGACCG	GATTATCTTC	CTCGACCAAG	GGAAAATCGC	TGAAGAAGGA	ACAGCTCAAG	23100
CCTTCTTTAC	CAATCCGCAA	ACCAAACGAG	CCCAAGGAATT	TTTAAACGTC	TTTGACTTTA	23160
GCCAATTCCG	CTCATATCTA	TAAAGGAGAT	TCTTATGAAA	CTATTCAAAC	CACTCTAAC	23220
TGTTTTAGCA	CTTGCCTTTC	CCCTTATCTT	TATCACTGCT	TGTAGCTAG	GTGGAAACGC	23280
TGGTTCATCC	TCTGGAAAAA	CAACTGCCAA	AGCTCGCACT	ATCGATGAAA	TCAAAAAAAG	23340
CGGTGAACTG	CAGATCGCCG	TGTTGGAGA	AAAAAAACCG	TTTGGCTACG	TTGACAATGAA	23400
TGGTCTTAC	CAAGGCTACG	CTACGATATT	GAACTAGGGA	ACCAACTAGC	TCAAGACCTT	23460
GGTGTCAAGG	TTAAATACAT	TTCAGTCGAT	GCTGCCAAC	GTGCGGAATA	CTTGATTTCA	23520
AACAAGGTAG	ATATTAACAT	TGCTAACTTT	ACAGTAACG	ACGAACGTAA	GAAACAAGTT	23580
GATTTGCC	TTCCATATAT	GAAAGTTCT	CTGGGTGTCG	TATCACCTAA	GAATGGTCTC	23640
ATTACAGACG	TCAAACAAC	TGAAGGTAAA	ACCTTAATTG	TCACAAAAGG	AACGACTGCT	23700
GAGACTTATT	TTGAAAAGAA	TCATCCAGAA	ATCAAACCTCC	AAAAATACGA	CCAATACAGT	23760
GACTCTTACC	AAGCTCTTCT	TGACGGACGT	GGAGATGCCT	TTTCAACTGA	CAATACGGAA	23820
GTTCTAGCTT	GGGGCCTTGA	AAATAAAGGA	TTTGAAGTAG	GAATTACTTC	CCTCGGTGAT	23880
CCCGATACCA	TTGGGGCAGC	AGTCAAAAAA	GGCAACCAAG	AATTGCTAGA	CTTCATCAAT	23940
AAAGATATTG	AAAAATTAGG	CAAGGAAAAC	TTCTTCCACA	AGGCCTATGA	AAAGACACTT	24000
CACCCAAACCT	ACGGTGACGC	TGCTAAAGCA	GATGACCTGG	TTGTTGAAGG	TGGAAAAGTT	24060
GATTAGTCAT	TAACCTCTAA	AAGGAACCTGG	ATTTTAAGCT	CCAATCCCTT	TTTAAGATT	24120
TACCTATAAC	ATCCTGAGTC	TATCTAACAT	GTTCAATCTG	AAACACAGTGT	ACATACTTTA	24180
TCTTCTATTG	CATATACATT	ATCACACATAAG	ATACGAATAT	CCTCTTCACT	ATGACTAGCA	24240
ATCAAAATTG	TTGTCCTTT	TTCACTAGAG	AGCTTTCTAA	ACAATGTTCT	CATATTTCT	24300
ACACTTGATT	TATCCAAGGC	ATTCATAGGT	TCATCTAGTA	AAAGAATAGA	GGGATTCTCC	24360
ATAATTGCTT	GAGCAATCCC	TAGCTTTTC	CTCATACCTA	GCGAATAAGT	TTTAACCTTC	24420
TGGTCTTTT	GCTCATATAG	ACCAACTATT	TTCACTGCTAT	CATTGATTTC	CTGATTACCA	24480
ACTACTCCTC	GTATGCTTGC	CAAATATTGT	AAATTCTAA	AGCCACTATA	ATAATTATA	24540
AAACCAGGTT	CTTCAATCAA	AGCTCCAAA	TTAGCTGGAA	TTTTCTCTC	AGGAACAATA	24600

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TTTCCCCAT TGATTAACAC TTCTCCATAA GACGGACTAT ATAAACCAGC TATTAATTAA	24660
AACAATACAC TTTTCCCTGA GCCATTCGCA CCAGTAATTC CTATAATTTC CCCCTGTTA	24720
CAACTAAAGT TAAGGTTTG AAAAACACAT GTCTTTTTA ATTCAACTC AATATTTTT	24780
AATGTAATTA TTTCATTCA TCTATAAACCC TCCTCTTTG ACGAGTGAAA TAGAAAATGC	24840
TTTGAAAAAG AAAGACTAA AATAGCAACT GAAGAAATAA ATCTCGTCCT ATATCTCCAT	24900
TCCCTCGATT CAAAATATAA AATAGATAAT TAGTCGATT TCCTACAAAT AGACCACCAA	24960
ACACAATCAT GAGTAAAAG AAACTAACGC AAGCAAAGTT CG	25002

(2) INFORMATION FOR SEQ ID NO: 49:

- (i) SEQUENCE CHARACTERISTICS:
- (A) LENGTH: 11443 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 49:

CAGGTACGGT GAGGCCAAC TAAAATATAA TTTTCATCTT GATTAGGAAT TTTATCAGTA	60
TTATGATAGT GAGCATTGCC ATTGATGGAC CATAAGAGCA ATACAACCAA TCCACGCAA	120
TAAGTATAAA ACATCCGATC TCCTTCGATT GTTTCTTGT TATTATTATA CCTTATCAA	180
GGAGGGCTGG CAAACTTTTC CCTTGACTAG ATACATATTT AGGATGAAAT TAGAATTCTG	240
TTAAAAAAA TGATATAATA GAATTATGG ATAAAATAA GATTATGGG TTAACCCAAA	300
GAGAAGTCAGG GGAAAGACAG GCTGAGGGTT TGGTCAATGA CTTTACCGCA TCAGCCAGTA	360
CCAGCACCTG GCAAATCGTT AAACGAAATG TCTTTACCCCT TTTAACGCT TTGAACTTTG	420
CCATTGCTTT GGCTCTTGCC TTTGTGCAGG CTTGGAGCAA TCTGGTCTTC TTTGCTGTTA	480
TCTGCTTTAA CGCTTTTCTT GGGATTGTGA CCGAGCTACG AGCCAAACAC ATGGTGGACA	540
AGCTCAATCT CATGACCAAG GAAAAGGTCA AAACCATCCG TGATGGTCAG GAAGTTGCTC	600
TTAATCCTGA AGAATTAGTG CTAGGAGATG TCATTCGTTT GTCTGCAGGA GAGCAGATTG	660
CTAGTGATGC CTTGGTTTG GAAGGTTTG CGGAAGTCAA TGAAGGCATG TTAACGGGAG	720
AAAGTGATTT GGTCCAAAAG GAAGTTGACG GCTTACTTTT GTCAGGAAGT TTCCTAGCCA	780
GTGGGTCACT TTTATCTCAA GTTCACCATG TCGGTGCAGA CAACTATGCT GCCAAACTCA	840
TGCTTGAGGC TAAGACCGTT AAACCCATCA ACTCCCGTAT CATGAAATCG CTGGACAAGT	900
TGGCTGGTTT TACTGGGAAG ATTATCATTC CCTTTGGTCT GGCTCTCTTG CTGGAAGCCT	960

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TGCTTTAAA AGGCCTGCCT CTCAAGTCAT CCCTTGTAAA CTCGTCGACA GCTCTTTGG	1020
GAATGTTGCC TAAGGAAATT GCCCTTTGA CCATTACTTC GCTCTTGACT GCAGTGATTA	1080
ACTTGGCCTT GAAAAGGTC TTGGTGCAGG AGATGTACTC TGTTGAGACC TTGGCGCGCG	1140
TGGATATGCT CTGCTGGAC AAGACGGTA CCATCACCCA AGGAAAGATG CAGGTGGAGG	1200
CTGTTCTTCC GTTGACGGAA ACCTATGGTG AAGAGGCTAT TGCCAGCATH TTGACTAGCT	1260
ACATGGCCCA TAGTGGAGGAT AAGAATCCAA CTGCCAAGC CATTGCCAG CGTTTTGTGG	1320
GAGATGTTGC TTATCCTATG ATTTCCAATC TCCCCTCTC GAGCGACCGC AAGTGGGGGG	1380
CTATGGAGTT AGAAGGCTTG GGGACAGTTT TCTTPAGGGC ACCTGAGATG TTGCTTGATT	1440
CTGAAGTCCC AGAAGCTAGG GAGGCCTTGG AGAGAGGATC ACGTGTCTTG GTCTTAGCTC	1500
TCAGTCAGGA GAAATTAGAC CATCACAAAC CACAGAAACC ATCTGATATT CAGGCTCTAG	1560
CCTTGCTGGA AATCTTGGAC CCCATTGAG AGGGAGCAGC AGAGACGCTG GACTATCTCC	1620
GTTCTCAGGA GGTGGGACTC AAGATTATCT CTGGTGACAA TCCAGTTACG GTGTCCAGCA	1680
TTGCCAGAA GGCTGGTTTT GCGGACTATC ACAGCTATGT AGATTGCTCA AAAATCACCG	1740
ATGAGGAATT GATGCCATG GCGGAGGAGA CAGCTATTTT CGGACGTGTT TCCCCTCATC	1800
AAAAGAAACT CATCATCCAA ACCTTGAAAA AAGCGGGACA TACAACGGCT ATGACAGGG	1860
ACGGGGTTAA TGATATCTTG GCCCTTCGTG AGGCGGATTG TTCTATCGTG ATGGCGGAGG	1920
GGGATCCAGC AACCCGTCAG ATTGCCAATC TCCCTCTCTT CAACTCAGAC TTTAATGATG	1980
TTCCTGAGAT TCTCTTCGAG GGTGCGCGCG TGGTCAATAA CATTGCCAAC ATCGCCCCGA	2040
TTTTCTTGAT AAAGACCATC TATTCTTCC TGTAGCAGT CATCTGTATT GCCAGTGCCT	2100
TACTAGGTG TGTAGAGTGG ATTTTGATTT TCCCCTTCAT TCCGATCCAG ATTACCATGA	2160
TTGACCAGTT TGTGGAAGGT TTCCCACCAT TCGTTCTGAC TTTTGAGCGA AATATCAAAC	2220
CTGTTGAGCA GAATTTCCTC AGAAAATCCA TGCTTCGTGC CCTACCAAGC GCTCTCATGG	2280
TCGTTCTCAG CGTCTGTGTT GTGAAAATGT TTGGCGCGAG TCAAGGTTGG TCTGAGTTAG	2340
AAATCTCAAC TCTACTCTAT TATCTCTTGG GGTCAATTGG TTTCTTATCC GTATTTAGAG	2400
CCTGCATGCC ATTACCCCTA TGGCGTGTCC TCTTGATTGT TTGGTCAGTA GGAGGTTTCC	2460
TAGCCACAGC TCTCTTCCCA AGAATTCAAA AACTGCTGA AATTCACCAAC TTAACAGAAC	2520
AAACCTTGCC TGTCTTATGGT GTCATGATGT TGGTCTTAC CGTGATTTTC ATCCCTGACCA	2580
GTCGTTACCA AGCGAAAAAA TAAATCAAAA CCACCGAGTGT GAACTGGTGG TTTGTTCTGC	2640
GGCTATAAGC CGCTTCTACC GGCCAGGGCC AAAGGCCAC CGAAATAGCT TCCTCGCGCA	2700
CCACTTTCCC GAGCAGGTGC TAAAGCACCT TAGTTACTTC CTCTTATTAA TTTCGCCAGT	2760

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AAACGGATCT ACTGACTCGA ATAACGTGAG CTGGTCTGCT ACTCTGTCTT CTTGTAATTG	2820
ATTCTGAATA TATTCAAGCTA TCACCTTCCTG ATTACGGCCT ACCGTATCTA CATAATAGCC	2880
TCTACACCAA AACTTGCATG TGCCATATTT GTATTTAAA TTGCGATGCT TATCAAAAT	2940
CATCAAACCTG CTCTGCCCT TTAAATAGCC CATAAAGGAC GAAACACTAA GTTTCGGAGG	3000
AATACTGATA AGCATGTGAA TATGGTCTGA ACAAGCATTG CTTTCATGGA TTATTACACC	3060
CTTACGCTCA CATAAGTCAC GTATGATTCT TCCGATACTA GCTTTGTATC TGCCATAAT	3120
GATTTGACGA CGATATTTGG GTGCAAAAC AATATGATAT TTACAATTCC ATGTGGTATG	3180
TGATAAACTT TGATTATCCT CTCTCATGAG GTACCTCCCTG TATGATATCT TGTAAGTGGCG	3240
GAGAAACAC TTCTATCTTA TCATTTTAGG AGGTTCTTT TGTTACCACG CTAAAGCTC	3300
TATGGAAcCA CTAGCATAGC TAGTGGTTTT CGGGAGACAA CAAGAAAGAC TGCAATCTGT	3360
GGATTGCGAGT TTTTATACG ATGGATCTAT CGTAGATCTG ATGTGCAAGG CCTACGTGCC	3420
GATCATCTAT CGGTGAACCC AAGAGCGACC CTCAGCCTG CTTGGATTGA GGTAATAGAT	3480
TCAAATATCT GTAGTTAGAC TATTGAAAGT TTGATGTAAG AAAGAGAAAG CGACAGATTG	3540
AAGTAATTTT AACTCTCTTC TATTGCTAGA ACAAATGGTC GGATAGGTTG GTAGTTGAA	3600
AATGAAGATG CTATCTATTG TAAATATGAA CATAGTGTAA TTTATTAGAA AATCGTTTG	3660
TTTATTTCTT ATCAAATACG AAAAGCACT TAAATATTTTCA AACTAAAATA GATGTTATGA	3720
AGAAAAAGTA AAATGATTTT GGCATAGTGA GGTTCTGTT TTTTGATAT CATATTTTG	3780
ATAAAAACAA AAATGTCCAT TGCAAAGGAC AAAATGCAGA GTATATTATT TTTTGAAAGC	3840
GATATAATGG ATTCACTAAAG GAGGTGTATC GTGTCTAGAA AACAGAACAA AATGGAAACG	3900
TTGTTGCTCC TTTTGCAGA TAGTAAGGAT TATATATCTG CTAAAGTATT GGGAGAAAAA	3960
TTAAATTGCT CTGATAAAAC GGTTTATCGC CTTGTCAAGG GAATCAACAA AGATTGTCCG	4020
GTAGAAGCAT TCATTTATC TGAAAAAGGC AGAGGTTCA AATTAAATCC AAGAAGTTCC	4080
CTCGTGGACG TTGATGGAA TTTTACAGAG GCTTTGATC CTGAAGTAAG GCCTGAAAAA	4140
TTACTAGAAC GTCTCTTGTGTT GACTGCTCCT AAGCCACATT CTATTTATGA TTTAGGAGAG	4200
GAATTCTACG TAAGCGAGTC AGTAGTACTA AAAGATCGTC AGATATTACA AGAGAGTCTA	4260
GCAATTATG GGTTAGATTT AAAATGAGA CAACGAAAGC TTTTTATTGA TGGGGATGAG	4320
GCTCAAATTC GTTCAGCCAT TCTAAATCTA CTGCCAATGT TTAATCAGTT GGATTTAGAG	4380
CAAATTACAC AGAATAAGGT TCAGCCTCTT GACGGAGAAC TTGCTCACTT TTGTTGGGA	4440
TTACTGATTA CACTTGAGAG AGAATTGGGG GTAAACATTG CCTATCCATA TAATATAAAT	4500

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ATTTTCCTCTC ACCTGTATAT TTTTATCAGT AGGAATCGTC GTAGTACTAG TATTCATGTT	4560
GTAGCACCTT CAAAACCTAC TATTGTTGAT GAGAAAATTT ACAGTGTCTG TCAAAAAATT	4620
ATTCAGAAGA TTGAAACAATA TTTTAGGATG AAGGTTGATG CAGTTGAGAT TGACTATCTT	4680
TATCAATACG TTGTATCTTC GAGATTGCAA AAACCATTTT CTTCCGGAA GCTTCCTTTT	4740
TCTCAGCGAG TTTTAGATGT CACTCATTAC TATTTTAGCC GTATGTGTAT GGACAATAGA	4800
GAGATTGAAA CGACAGATCC TGACTTTGTT GACTTGGCGA GTCATATCAG TCCCTTACTG	4860
AGGAGATTAG ATAATAGAGT ACAGATTAAG AATAGTCTTT TATCACAAAT TCTTTTAACC	4920
TATCCTAATC TGGTTAAAGA GTTAACAACT ATTCTAAAG AAGTGAGTCT AGTATTTGGT	4980
TTTGCTTCCT TGAGTCTGGA CGAGATTGGT TTTCTAGTCT TATATTTGC ACGGTTCAA	5040
GAAAAGCGAG CACGTCTCT AAAAACAGTA GTGATGTGTA CATCAGGTGT CGGAACCTCA	5100
GAGCTTTAC GAGCACGATT AGAAAAGCAA TTTCTGAAT TGGATATTAT TGATGTAGTT	5160
GCTTATCATC AATTAGATGA GCTGATAAAAT CTATATCCAG ATTTAGATTT CATTGTGACG	5220
ACGGTAGCTT TGCAGGAACC AGCAAGTGTG CCGTTTGTCC TAGTTAGTGT TTTCTAACCC	5280
GAGGGTGATA AACAAACGTCT TCAAGCAAA ATTCAAGGAGA TAAACTATGA ATAATCTTC	5340
GCTTGTCTT ATGGATATAT CTGTTAAAAA TCGTCAAGAA GCCTACAAAG AATTAGCAAA	5400
TCAAATCAGC CTTCTGTCTT CTGAAGATAC AGAAAAAATA GAAGAGCTTC TATATTACCG	5460
TGAGAGACAG GGAAGTATAG AGGTTGCTAA AGGTGTTCTT CTACCCACATT GTGAAACAAA	5520
CTTTCAACAT CATGTCTTAG TGATTACTAG ATTAACATCA CCTATCAGAG AATGGTCGAA	5580
GGATATCCAG TGTGTTGACC TTATTATCGG TTGGGCCATT GCAGTATCAC AGGACAAGTC	5640
ATGTATTTAA ACATTGTGA GAAGACTAGC AGATGAATCA TTCATAAAATC AATTAAAACA	5700
GTAAACAAA GAAGAATTAC GGGAGATAAT ATATGGAAT CAAAGATATT CTTAATGTGA	5760
GTCTGATCCA GACGGATTTA CAGATGCAGA GCAAAGAAGA GGTTTTGAG GCATTAGCTC	5820
AACTATTGGT TGAGACGGGT TATGTGTCTG ATAGAGACCA ATTTATCGAA GGTCTTTATC	5880
AGAGAGAGGC AGAAGGACAG ACCGGTATTG GGAATTATAT TGCTATTCCC CATAGCAAGA	5940
GTTCTGCTGT GGAGAAGGCG GGGGTAGTCA TAGCTATAAA TCACAATGAG ATTCTTGGG	6000
AGACCATTTGA TGGGAAAGGG GTCAAAGTAA TTGTACTCTT TGCAGTTGGT GATGATACAG	6060
AAGCTGCTAG GGAGCATTG AAGACCTTAT CACTCTTGC TCGAAAACCTT GGTAATGACG	6120
AAGTTGTTGC CAAATTAGTT CGGGCTCAGA CATCTGATGA TGTGATTGCA GCTTTTTGTT	6180
AATAAGAAA AATTGGAG GGTATCCGTA TGAAAATTGT TGGTGTGCA GCTTGTACTG	6240
TGGGAATTGC CCACACTTAT ATTGCACAGG AAAAATTAGA GAATGCCGCA AAGGTAGCTG	6300

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GACATGTGAT TCATGTTGAG ACTCAGGGGA CAATAGGGT AGAAAATGAA TTGAGTCAG	6360
AGCAGATTGA TGCAGCGGAT GTAGTTATTT TAGCAGTTGA TGTTAAGATT TCTGGTATGG	6420
AACGCTTGA GGGTAAAAAG ATTATCAAGG TTCCAACAGA AGTGGCAGTC AAATCTCCCA	6480
ATAAAATGAT TGCTAAAGCT GTTGAGATTG TTACGAAATA ACTGAAAATA TTTAAGGAGA	6540
AAATATATGT TGAAACACTT AAACTTAAAAA GGTCACTTAT TGACAGCCAT TTCCATATATG	6600
ATTCCAATTG TTTGTGGTGC AGGATTCTTA GTTGCCATTG GTTPAGCAAT GGGGGGTGGT	6660
GTTCCGTGACG CTCTTGTAGC AGGAAAATTC ACTATCTGGG ATGCTTGTAGC AACTATGGGT	6720
GGTAAAGCCC TTGGTCTCTT GCCAGTTGTT ATTGCTACAG GTTGTCTTA CTCGATTGCT	6780
GGTAAGCCAG GGATTGCACC AGGTTTTGTT GTTGGTCTAA TTGCCAATTC TGTTGGTTCA	6840
GGGTTATCG GTGGTATCTT GGGAGGTTAT ATAGCTGGTT TCTTGGTTCA AGCGATTATT	6900
AAAAAGGTCA AAGTACCAAA CTGGATTAAA GTTAAATGC CAACCTTGAT TATTCCCTTT	6960
GTAGCCTCTT TGGTAAGTAG TTTGATTATG ATTTATATTA TTGGAGCGCC TATCGCAGCC	7020
TTTACCAACT GGTTGACGAG CTTATTACAA AGCTTGGAA GTGCTCAA TGTTTGATG	7080
GGGGCAGTTA TTGGAATTCT CAGTGCTGTT GACTTTGGTG GCCCACTTAA TAAAACAGTC	7140
TATGCCCTTG TGTTGACTTT ACAGGCTGAA GGTGTGAAAG AACATTGAC TGCTTACAA	7200
TTGGTGAATA CTGCTACACC ACTTGGATTT GGATTGGCT ATTTTATGCC GAAATTACTC	7260
AAAAAAAAATA TCTATACTCA AGAGGAAATC GAAACATTGA AATCGGCTGT TCCTATGGG	7320
ATTGTCAATA TTGTTGAAGG TGTAATTCCG ATTGTTATGA ATAACCTGGT TCCAGGTCTC	7380
ATTGCAACAG GTATCGGTGG TGCTGTTGGT GGTGCTGTTT CTTTGACAAT GGGTGCTGAT	7440
TCTGCTGTGC CATTGGTGG AGTGCTTATG TTACCAACCA TGACTCGTCC AGTAGCTGGT	7500
ATTTGTGCCT TGTTAGCTAA CATTGTAGTC ACAGGACTTG TCTACGCGAT TTTGAAAAAA	7560
CCAATAAAAC ATGCAGAACC AGTTATGACT GTTGAAGAAG AGATTGATTT GTCAGATATT	7620
GAAATTGGTGT AAGAGGGTAA CGATGTCAAG AATTGAATT TCACCATCTT TGATGACCAT	7680
GGATTGGAC AAATTCAAAG ACCAGATTAC TTTTTGAAAT GATAAAGTAG CATCTTATCA	7740
TATCGATATT ATGGATGGCC ATTTGTTCC CAATATTACC TTGTCCTT GGTTCATTC	7800
AGAAGTTCAA AAAATTAGTG ACACACCTTT ATCAGTTCAT CTGATGGTCA CAGACCCAAC	7860
CTTTGGGTA GATCAAGTTC TCGATTTACA ATGTGAGTAT ATTTGTTATTC ATGCTGAAGT	7920
TCTGAATGGT CTTGCTTTTC GTTGATTGA TAAAATTCAAT GATGCAGGTC TAAAGGCTGG	7980
TGTTGTCCTT AATCCTGAAAC CACCTGTTTC TACAATCTTT CCCTACATTG ATTTACTTGA	8040

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CAAAGCAACT ATTATGACTG TAGATCCAGG TTTTGCAGGA CAACGCTTTT TGGAGTCTAC	8100
CTTGTATAAA ATCCAAGAAC TCCGTCAGCT TAGAGTTCAAG AATGGTTATC ACTACATCAT	8160
TGAGATGGAT GGTTCTTCGA GTCGTAAGAC TTTCAAACAA ATTGTATGG CAGGACCAGA	8220
TATTTATGTT ATAGGTCGCA GTGGATTATT TGGTTGGAT GACGATATTG CCAAAGCTG	8280
GGATATCTGT TCTAGAGATT ACCAAGAAAT GACCGGAAAA ACAATGCCAA TCAAATAATG	8340
GTTTGAGAAG AAATTTATTA GTTAGGAGGA ATATATGTCA CTACAATCAG TTAACGCCAT	8400
TCGTTTCCTT GGCGTAGATG CTATTAACAA ATCTAATTCT GGTACCCGG GAATTGTAT	8460
GGGTGCTCGG CCAATGGCTT ATAGCCTATT TACAAAGCAC CTTAGAATTAA CACCTGAGCA	8520
GCCAAACTGG ATTAACCGAG ATCGCTTTAT CTTGTCGCG GGTACATGGAT CAATGCTACT	8580
GTATGCTCTC TTGCATTAA CAGGGTATAA GGATGTATCC ATGGACGAGA TTAAAAATT	8640
CCGGCAATGG GGATCTAAGA CACCTGGTCA TCCCTGAAGTG ACGCATACTG CTGGTGTGGA	8700
TGGCACATCT GGTCCGCTTG GTCAGGGAT TTCTACTGCC GTGGTTTCC CCCAAGCAGA	8760
GCGTTTTTA GCTGCTAAAGT ACAACAAAGA TGTTTCCCT ATTTTGACCC ATTATACTTA	8820
TGTTATCGCT GGAGACGGTG ACTTCATGGA AGGAGTGTCT CGGGAGGCGG CTTCTTATGC	8880
AGGTCATCAA GCTTTAGATA AGCTTATCGT CCTCTACGAC TCCAACGACA TCTGCTTGG	8940
TGGTGAGACC AAAGATACTT TCTCTGAAA TGTTCGCGTC CGTTACGATG CTTATGGTTG	9000
GCATACAGTT CTGGTAGAAG ATGGAACAGA TTTAGCAGCA ATTTCTACAG CAAATGACAC	9060
GGCCAAGTTT TCTGGTAAAC CGAGTTTGAT TGAAGTGGAAA ACGGTAAATTG GTTACGGCTC	9120
ACCCAATAAA AGTGGTACAA ATGCTGTTCA TGGTGCACCA CTAGGAGCAG AAGAAACAGG	9180
AGCAACTCGT AAGTTTTGG GATGGGATTA CGATCCATTG GAAGTACCGAG AGGAAGTATA	9240
TTCTGATTTA AAGACAAATG TAGCGGATCG TGGTCAGGAG GCATACGATG CTTGGGCTAG	9300
TTTGGTGTCT GATTACAAGG TTGCTTATCC CGAAGTTGCT AGTGAGATTG ACGCTATTGT	9360
AGCTGGAAA TCCCCGTAA CCATTACTGA AAAAGACTTC CCTGTCTATG AGAATGGCTT	9420
CTCTCAAGCA ACTCGTAATT CGTCCCAAGA TGCTATTAAT ACAGCAGCAG TTTTACCAAC	9480
CTTCTTAGGT GGATCGGCAG ACTTAGCTCA CTCTAACATG ACCTACATCA AGGCAGATGG	9540
CTTPACAAGAT AAATATAATC CATTAAACCG CAATATTCAAG TTTGGGTAC GTGAATTTC	9600
CATGGGAACA ATCCCTCAATG GAATGGCTCT TCATGGTGGT TTACGAGTTT ATGGCGGAAC	9660
CTTCTTTGTT TTCTCTGACT ACGTCAAAGC TGCTATTCCG CTATCAGCCA TTCAGGAGTT	9720
GCCTGTAAC TATGTCTTTA CCCATGATTC AATTGCCGTT GGTGAAGATG GTCCAACCTCA	9780
TGAACCAGTT GAACATTGG CAGGTTACG CTCAATGCCA AACTTGACTG TTATCCGTCC	9840

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AGCGGATGCC CGTGAAACTC AAGCGGCTTG GCATCATGCC TTGACCAGTA CCACCACTCC	9900
AACTGTCATT GTCTTAACCC GTCAAAAACCTT GGTAGTTGAA GAAGGGACAG ACTTTGGTAA	9960
GGTCGCTAAA GGAGCCTACG TCGTGTATGA TACCCCGGGA TTTGATACTA TTATCATTGC	10020
TACAGGATCT GAGGTCAATC TAGCTATCAA AGCTGCTAAG GAATTGGTTT TACAAGGTGG	10080
TAAAGTACGT GTGGTATCTA TGCCCTCAAC CGAACTATTT GATGCTCAAG ATGCTACCTA	10140
CAAGGAAGAC ATTTTACCAT CTAAGACTCG TCGTCGTGTG GCCATTGAAA TGGCAGCGAC	10200
CCAAAGTTGG TACAAGTATG TTGGTTTGGG TGGCGCGGTC ATCGGTATTG ACATCTTCGG	10260
TGGCTCTGCC CCAGCTCAGA CTGTGATTGA TAATTATGGA TTTACGGTAG AGAATATCGT	10320
TGCTCAAGTT AAGTCCCTAT AGAAAACCAAT TACAATGAAG ATACAGCTGT TGTCAGACTA	10380
GCAGATGTAG TGATAGACAC TAATCAGATG ATTGGTTATT TAAAAACTGT AATGAAAATG	10440
TAATAATTAA TCTACGAAAG TTATAGTACA TAGTATAACAC AATAGAGTAT ACCCTGAAAC	10500
GGTTGCGAAG TACGCTAAC TACCTTGCTAC TGATCTAGAT AGTTTCTTTA ATCAATAAAC	10560
ACAGCATCCA CAGATTGACT TAGGATATTG TAAGTTTTT GAAAGCTAGA GAGAAGGCT	10620
CTAAAATTAA AAAACGCATA GTATAGGATG TTGAAATGAT GAACTGCACC CCAAAAGTTA	10680
GACAGAAAAA AATCTAACTT TTGGGGTCTT TTATTTATGA AATTAACCTA TGATGATAAA	10740
GTTCACTCT ATGAACCTAG AAAACAAGGA TATATCTTAG AGAAGCTTTC AAATAAATT	10800
GGGATAAAATA ATTCTAATCT TAGGTACATG ATTAAATTGA TTGATCGTTA CGGAATAGAG	10860
TTCGTCAAAA AAGGGAAAAA TCGTTACTAT TCTCCTGATT TAAAACAAGA AATGATTCT	10920
AAAGTCTGAC ATGAAGGCTG GACTAAAGAT AGAGTTCTC TTGAATACGG TCTCCCAAGT	10980
CGTACGATAC TTCTTAACCTG GCTAGCACAA TACAGGAAAA ACGGGTATAC TATTGTTGAG	11040
AAAACAAAAG GGAGAGTACC TGAGAGCGGA GAATGCCATC CTAAAAAAGT TAAGAGAACT	11100
CCGATTGAAG GAGGAAAAAG AGAAATAAGA AAGACAGAAA TTGTTCAAGA ATTAATGACT	11160
GAGTTTCGTT TAGATCTTCT TCTAAAAGCC ATTAAACTAG CTCGTTGGAC CTACTACTAT	11220
CACTGAAAC ACCTAGATAA ACCAGATAAG GACCAAGAGC TTAAAGCTGA AATTCAATCC	11280
ATCTTATCG AACACAAGGG AGATTATGCT TATCGCCGGG TTCATTTAGA ACTAAGAAAT	11340
CGTGCTTATC TGGTAAATCA TAAAAGAGTT CAAGGCTTGA TGAAAGTACT CAATTACAA	11400
GCTAGAATGC GACAGAACG AAAATATTCT TCTCATAAAG GAG	11443

(2) INFORMATION FOR SEQ ID NO: 50:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 5338 base pairs

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- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 50:

CCAATTACAT TATATTATCA AAATCGTCGA AACTGGCTCC ATGAATGAGG CAGCCAAGCA	60
ACTCTTTATC ACTCAGCCAA GTCCTCTCCAA TGCAGTGCAGA GATTGGAAA ATGAAATGGG	120
CATTGAGATC TTTATCCGCA ATCCAAGGG AATCACCTTG ACCCGTGATG GCATGGAGTT	180
TCTCTCTTAT GCCCCGTCAGG TTGTCGAGCA GACCCAGCTT CTGGAGGAAC GCTATAAAAA	240
TCCTGTCGCC CACCGCGAAC TCTTTAGCGT TTCTGTCAA CACTATGCCT TTGTGGTCAA	300
TGCCTTTGTC TCTTGCTCA AGAAAAGCGA TATGGAGAAA TAGGAACCTCT TCCTTCGTGA	360
AACTCCGACT TGGGAGATTA TCGACGACGT CAAGAACTTC CGCAGTGAGG TCGGGGTCTT	420
CTTCTTAAAC AGTTACAACC GTGATGTTTT AACCAAGATG CTGGATGACA ATCACCTGCT	480
AGCCCACCAT CTCTTCACAG CGCAACCGCA TATCTTGTC AGCAAGACCA ACCCTCTGGC	540
AAAGAAAGAC AAGGTGAAAC TGTCTGATTT GGAGAATTTC CCTTACCTCA GCTATGACCA	600
AGGGACGCAC AACTCCTTCT ACTTTTCAGA AGAGATTCTT TCTCAAGAAC ACCACAAGAA	660
ATCCATTGTC GTCAGTGACG GTGCCACCCCT CTTTAATCTC TTGATTGGTT TGGATGGTTA	720
TACCATGGCG ACAGGGATTT TGAACAGCAA CCTAAACGGA GACAATATCG TTTCTATCCC	780
ACTGGATATT GATGACCGCA TCGAGCTGGT CTATATCCAG CATGAGAAAA CCAGCCTATC	840
TAAGATGGGC GAACCGCTTA TAGACTATCT CCTAGAAGAA GTTCAGTTG ATAGTTGAGA	900
AATGATAAGA ACCAATATGT AGGCTAGCAA CAACCTGCAC ATTGGTTCTT TTTACTTATA	960
ATTAAAAGTT TCCCCGCCA ACTTATCAGC TAGCTGGGA AAGAGAGTAT AAAACTTATG	1020
GGCTAGGTTA AACAAAATCG GGAGATTGAG TTCTCGTTG TTTTTCCCTA TAATCTTGAC	1080
AATCTTTTA GCCACTGCAT CTGGTTCTAG CAGGAAGCGA TCAACCGATT TAAGATAAGT	1140
TCCATCTGGG TCGGCTTGTT CGAAAAATCC TGACCGATT GGTCTGGAT TGACTGTTGT	1200
CACATAGACT CCATAGGGCA TAAGTCGAG TCGCAGAGCA TTTGAAAAAC CAATAGCCGC	1260
AAACTGGTC GCTGAGTAAA GACTAGACTT GCCAGTAGCT ATTAGACCTG CCATGCTGAC	1320
GATGTTGATG ATATGCCCTT TGCTGCTTTC CTTCATACGA GCCGCAAGGT GACGAGACAG	1380
ATTCAATCAGG GCAAAGGTAT TGACCTCAA CATCTGGTGA ATATCTTTAT CAGCAATCTG	1440
GTCAAATCCC TCAAAAATCC CGTAACCAGC GTTGTAAATC AAGACATCAA TCTTGCCATA	1500
CGGGAGATAA AGATCAGTTA CCAGAGCTTC TAGGGCTGAA TCGTCGGTAA TATCAATTTC	1560

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AATCAATTCT GCATGGGAAT AATTTCCGTA GAGTTGGGCT AATTTTCCT TATTTCTACC	1620
AAGCAAGATG AGTTGGTCAT TGCGCAGGAG TTTGACCATT TCTTGAGCTA GACCACCGCT	1680
AGCTCCGCTA ATGAGAAATAG TAGGCATACT TATCCTTCT GTGACTGCTA GATTTCCACT	1740
TCTTCCAAGT CTTTGACAC ATGGACATTT TCAAAAATTG TGGCAGCGTC TTTCTTGAGT	1800
TTGCTAATAT CTTTGAGAG GAAACGGGCA CTGATATGGT TGAGTAGGAG GCGTTGGCA	1860
CCTGCTTCTA CCGCTACTTG TGCGAGCTTGC ATATTAGTTC AGTGACCATG GTTACGAGCA	1920
ATTTTTTCAT CACCCCTGCC ATAAGTGGAC TCATGAACTA GGACATCTGC ATTGACAGCC	1980
AGACGCACAC TGGCACCCGT TTTTCGAGTG TCTCCTAAA TAGTGATAAT CTTACCTGGA	2040
CGTGGCGCTG AGATATAGTC TGCGCTTGC ATTTCAAGTTC CGTCTCCAA ACAAGATCC	2100
TGGCCGTTTT TGATTTTACCA AAAAAGCGGG CCGAACGGAA CACCAAGCAGC CTTGAGTTT	2160
TCAGCATCCA GCGTCCCTTC TAGATCCTTT TGCGATGACAC GATAGCCAAC ACAGAAAATA	2220
GTGTGGTCCA GCTCCTCTGC ATACACAGTG AATTTATCGG TTTCAAGAAT TTTACCCAGA	2280
GAATCTTGGT CAAACTCATG GAAATGAATG CGGTAGGGCA GACGAGAAC TGACACACGA	2340
AGGCTGGTTA AGACAAATGA CTTGATTCCCT TGAGGTCCGT AGATTTCCAA ATCTGTCTGC	2400
TCTTCATTGG CCTGAAAGGC ACGGCTAGAA AGGAAACCTG GCAAACCAAA AATGTGGTCT	2460
CCATGCAGAT GGCTAATAAA GATTTTGCTG ACCTTACGTG GTGCAATTGT GGTTTCCAGA	2520
ATGGGATTTT GCGTACCTTC TCCACAGTC AAGAGCCAA CTTCGTTAAT CTCATCCAA	2580
AGTTTCAGGG CGAGACTTGA AACGTTGCGG GCTTTAGAGG GCTGACCAGC CCCCGTTCT	2640
AAAAATTGAA TATCCATTG ATACTTCTA ATTAATCAAT ATATAACATG GCTGTGCGGT	2700
TTTCCGATCG GAAATAGCGT TTGCCAGAAA AAGCAGCAGC TTCTGCAAT AAATCCTCTT	2760
GGCTGTAGCC TTTGAGACGT TTTGACCAT CAGCCAATCT TTCCAAATCA GTCAAAGCTG	2820
TGAGACTTTC TAGGCTGATA ACTTCCTCGT CCTCGACAGG CTTCATGTAA ATCTTACCAAG	2880
ACTCTCAAA GACTAATTGA TGGGGAAAA TTGCGCAAT TTCAAAGAGC AAGTCATCCG	2940
AGATTTCTC CTCATTTCA AAGAAAATCC GACCAAGGCC GTCACTCTCA TAACAAAAC	3000
CAAAGGATT ACCAGACAGA TTAAGCCGAA TAAAAGGCTT ATTTTCTAGG GTGAAACTTG	3060
GCTCAGTATT GTAAAGATTG AGTTCTGAC TGAGTTCTGC AAAATAATCC GTCGCAGCCT	3120
GAGGACTCTT TTTCTGATAG AGTTCTGCAA AGTAGGCATT AACACACATT GGCGGAGGTG	3180
TAATAAGTGT TAACTGCTCC TGATCTGTTT TACCAAGCTAG AAGCTGATCC AGATAGACCT	3240
TGTCCAGACT TGTATAACCT CCATACTTTA GAGCCAAAGT TTTAATATCA GTCATAAAAT	3300

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TCTTCTAAC	3360
TCCATTATT	
TTTCTCGGAA	
ATGTAGCCTG	
TAATCACTTC	
GCCGTCTTCC	
TGATAATCAC	3420
GTTCTCCAG	
AATTGCAACA	
CTCTCTAAAT	
CATGAATCTT	
GTAGGACTTT	
GAAAAAGGCA	3480
CTCGCAGGGT	
AAATGCTTCA	
AAAATTCCCT	
TAATCTTATC	
TAGCAATAAT	
GCTTGCAAGT	3540
TTTCACGACT	
GTCCCTCAGAC	
TTGGCAGAAA	
TGAGGGTATA	
TGGCGTTGG	
GTAGGGCTGA	3600
AATCCTCCAC	
CAAATCCGCT	
TTATTATAAA	
GCGTCAAGTG	
AGGAATATCT	
TCCATGTCCA	3660
GGTCTTCAT	
GATGGAGAGA	
ACCGTTTTT	
CATGCTCCTC	
GTGGTAAGGA	
TTGCTAGCAT	3720
CGATAACATG	
AACCAGAAGG	
TCCACATGCT	
TGCTTCTTC	
CAAGGTTGAC	
TTGAAAATGG	3780
ACACCAACTC	
TGTCGGCAA	
TCTTGGATAA	
AGCCAACGGT	
ATCTGTCAAA	
GTTACTTGG	3840
GATTCCTCC	
CAGATGAATA	
CTCTTGGTTG	
TCGCATCCAG	
AGTCGCAAAG	
AGCTCATCTG	3900
CTTCATACTG	
GGTCTTACTG	
GTCAAGATGT	
TCATGATAGT	
TGATTTCCCA	
GCATTAGTAT	3960
AACCAATCAA	
ACCAATCTTA	
AAAGTGCTAG	
ACTCCAAACG	
TTTTTCTCTG	
ACAGTCGCAC	4020
GATTTTCTC	
AACCACCTTG	
AGCTGGCGCT	
CGATATCCGT	
GATTTGATTG	
CGAACGCTAC	4080
GACGGTTAG	
CTCCAGCTGG	
CTTCACCAAG	
GACCACGGGA	
ACCAATTCCC	
CCTgCCTGAC	4140
GGCTGAGCAT	
AATCCCCTGA	
CCAACCAAGC	
GAGGCAAAAG	
GTATTTGAGT	
TGGGCTAGGT	4200
GGACTTGGAG	
CTTCCCTTCA	
TGGCTTCAG	
CCCGCATGGC	
AAAGATATCC	
AAAATCAACT	4260
GCATACGGTC	
AATGACCTTA	
ACACCGAGAA	
CTTCCTCTAG	
ATTGACATTC	
TGCCCTGGGG	4320
TCAGACGATT	
GTTGACGATG	
ACAGTAGTGA	
TTTCTTCTGC	
ATCCACCATA	
AGCGCAATCT	4380
CTTCCAACCT	
ACCAAGGCCG	
ACGAAGGTCT	
TGGAATCATA	
TTTTTCACGT	
TTTTGTCTGT	4440
AGCTATCTAC	
AACGACTGCC	
CCTGCCGT	
TCGCTAAACT	
AGCCAATTCT	
TCCATGGGAGA	4500
GGTCAAAACT	
GTCCATACCC	
TGCAATTCCA	
CACCAATCAG	
CAGGACTCGC	
TTGGCAAAAC	4560
GACGAGTCGC	
CTGTTTAAGA	
CTCTCACTAG	
CTTCCTCCAA	
GGTCTGCTCT	4620
CCACGGAAAT	
AAGGAAAGAG	
TTCCATTATAG	
CCAATTCCCT	
TAGCAGCCTG	
TACATTAGGG	4680
GAATGGTCAA	
ACAGCCACTT	
GGCCTCATCC	
AAAAGCCCAG	
CCTCAAACAT	
CAAATCCACT	4740
CGGTGGTTGA	
TACGCTCAT	
AAGTTGACTA	
CGTTCATCAT	
CCAAGCGAGT	
AATCAGCGGT	4800
TCATACAAAGG	
TCTCTTGATT	
TTCCAAATCC	
TGACCAAAAT	
GGGCAATTTC	
TAAGGCACGC	4860
ATAGCACGAC	
GACGATTAAA	
CTGGGAAATC	
TCAAGGCCTG	
CTTGATCCAC	
CAAATGGGCT	4920
AATTCCCTCAT	
CTGAATATGG	
CTCCAAACTA	
GCTCGATAAG	
CTAAATCTC	
CTCATGAGGA	4980
GTCTCCCCAC	
CTAGGTGGTA	
ACCTTCTAGC	
AAGCTCTGGA	
TATAAAGTCC	
	5040
	5100

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AGTCCCACCG GCGATAATGG CTAGCTGCC ACGGTTGTGA ATACCCCAA TAGTCATCTT	5160
AGCTTCTGAA ACAAAATCAA AAGCCGAGTA AGACTCGGTT ATCTCTCTAA CATCGATTAA	5220
ATGATGAGGA ACAGCTGCCT GCTCTTCTGG ACTAGCCTTG GCCGTCCCAA TATCAAGTCC	5280
TCGATAGACT TGCTGGCTAT CTCCACTAAC CACTTCGCCA TTAAAACGCT TTGCGGGG	5338

(2) INFORMATION FOR SEQ ID NO: 51:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 19446 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 51:

CGGAAACCCA TCTAGTCTCC ATCGTTGGG AGACCAAGCA ACACGAATCT TAGATGCTTC	60
TCGCCAACAG ATTGCAGATT TAATCGGTAA GAAAAGCGAT GAAATCTTCT TTACCTCGGG	120
TGGAACAGAA GGGGATAACT GGCTTATCAA GGGTGTGGCC TTTGAAAAAG CTCAGTTTGG	180
CAAGCACATC ATTGTTTCAG CCATTGAACA TCCAGCAGTC AAAGAGTCAG CCCTCTGGTT	240
GAAAAGTCAA GGATTGAAAG TGGATTTGC TCCAGTTGAT AAGAAAGGCT TGGTCGATGT	300
TGAGGGTTA CAGGTTTGAT ACGGCATGAT ACAATCCTCG TTTCCATCAT GGCTGTGAAC	360
AATGAAATCG GCTCTATCCA ACCTATTGAG GCTATTTCAG AATTCTTGGC AGACAAGCCG	420
ACTATTTCTT TCCACGTTGA TGCGGTTCAAG CGCCTTGCCA AAATTCCGAC TGAAAAGTAT	480
CTGACAGAAC GGGTGGATTG CGCGACTTTC TCTAGTCACA AGTTCCACGG GGTTGAGGT	540
GTTGGCTTTG TCTATATCAA ATCTGGCAAG AAGATTACAC CTCTTCTTAC AGGTGGTGGC	600
CAGGAGCGAG ATTATCGTTC GACAACGTAA AATGTGGCAG GGATPGCAGC GACAGCCAAG	660
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CACGCCCTTG AAGACTATGA TATTTTCATC TCAACAAACCT CAGCTTGTTC ATCTAAGGCA	900
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CCCTTTTGG TCAGTATCGA CACGAGACTC AACGTAAACG ACTTCACGAA TGACATCCTG	14460
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AAGACCATCA CTAACAAGTT GGTCGATAAG AGTATTTCT TTTTGGTGC GAGTATTGAG	14580
TAACTGATAG AGATTTCAA TCAAGTCACC ATATATAATG GAAAATCCAG TTTCTTTACG	14640
AAAAACGTCA CTATCTCGA AGTCAACCAA ATAAGAAAAG CCTAAAGTT GAAAAGCAAC	14700
AGTATAAAA ATATCTGCTC TCAGTTCATC TTCTGATTGA AAAATGTCA GCAGGTCTGT	14760
TTTTTTATCA GCTGCTAGGA TAGAAAGTGG GTAGTTGGTG TCTTGATAAG TGAAAAGAA	14820
ACGACGTAAA AAGGTTCAA GTGAGTCTTT GTGATTGGCT GTATTTGTAA AATCAAAGCC	14880
ACATTTTTT AGTTCAAGATA AGACATTTTC TTTGGAAAA TTGATATAAC TATATTGATT	14940
AAAACGCATA GAACCTCCAT ATAGAATGAC ACTTAAGGTT ATTATATCAA AAAAAAAGCA	15000
GAAAGGAAT TCTTAACCTTC AAAAGGAAT AATCCAATAA AAATGAATAA AGTACTAAAT	15060
TCAATATAGA GAACAGAGTA ACAATAAGAA TAAATAGATA GGGTATAAAA GTTCTAGGAG	15120
ATTTATATTA TATGCTTTCT ATTTTTATAT ACAATATAGT ATAAATATAA AAATGATGAC	15180
AAAAATACAA ATGAATAGAA AATAAATTAG TAAGCTGATG AAATTTTCT CAAGAGAAGC	15240
CATTTATAGG TGAAAATGGT ATAATATAGT GAGAAGGATA GAGGAGAAGT GTAAATTGAT	15300

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CGCACAACTA GATACAAAAA CAGTCTATAG TTTTATGGAA AGCGTCATTT CGATCGAAAA	15360
GTATGTGAGA GCAGCTAACG AATAACGGCTA CACTCATTG GCTATGATGG ATATTGACAA	15420
TCTTTATGGC GCTTCGACT TTCTAGAGAT TACAAAAAAA TACGGCATTC ATCCTTGCT	15480
AGGGCTGAA ATGACAGTGT TTGTAGATGA TCAGGGAGTG AATTTCGCT TTTTAGCTCT	15540
ATCTAGTGTG GGCTATCAGC AGTTGATGAA CCTTTCGACA GCCAAGATGC AGGGGGAGAA	15600
AACTTGGTCA GTCCTGTCAG AGTACCTGGA GGATATCGCG GTCATTGTGC CTTATTTGAA	15660
TAGAGTTGAG TCGTTAGAAC TAGGCTGTGA TTACTATATA GGGGTTTATC CAGAAACACT	15720
AGCAAGCGAA TTTCATCATC CTATCTTACC TCTTTATCGG GTCAACCGTT TTGAAAGCAG	15780
GGATAGAGAA GTTCTTCAG TTTAACAGC GATTAAAGAA AATCTACCGC TCAGAGAACT	15840
TCCCTTGCCT TCGAGACAAG ATGTCCTTAT ATCAGCAAGT TCTTTAGAGA AACTATTCCA	15900
AGAGCGTTT CCGCAAGCTT TGGACAATT AGAAAAGCTT ATTCAGGCA TTTCTTACCA	15960
CTTGGATACT AGTCTGAAAC TGCCTCGTT TAATCCAGCT AGACCAGCAG TAGAGGAGTT	16020
GAGAGAGCGT GCTGAACCTGG GGCTTGTCA GAAGGGGTTG ACTAGTAAAG AATATCAAGA	16080
TAGACTAGAC CAAGAATTGT CTGTTATTCA TGATATGGC TTTGATGATT ATTTCTTGGT	16140
TGTTTGGGAT TTGTTGCCCTT TTGGACAATC GAATGGCTAT TATATGGAA TGGGAAGGGG	16200
TTCTGCAGTA GGCAGTTGG TTTCTTATGC CTTAGACATC ACAGGGGATTG ACCCAGTAGA	16260
AAAAAACTG ATTTTGAAAC GCTTTCTTAA TCTGAAACCC TATACCATCC CTGATATTGA	16320
TATTGATATC CCAGATATTT ATCGTCCAGA TTTTATCAGA TATGTTGGTA ATAAATATGG	16380
TAGTAAACAT GCGGCACAAA TCGTTACTTT TCAACCTTT GGAGCCAAGC AACCTCTTCG	16440
AGATGTCTTG AAACGCTTG TGTCGCCAGA GTATGAATTA TCTGCAATTA CTAAGAAAAT	16500
CAGTTTCGT GACAATCTTA AGTCGGCCTA TGAGGGAAAT CTCCAGTTTC GTCAGCAAAT	16560
CAATAGTAAG TTAGAATACC AAAAGCTTT TGAGATTGCT TGCAAGATAG AGGGCTATCC	16620
AAGGCAAACC TCTGTCCATG CGCCTGGTGT TGTAATTAGT GACCAAGATT TAACCAACTA	16680
CATTCCCTCTA AAGTATGGTG ATGAAATTCC ACTGACTCAG TATGATGCTC ATGGAGTTGA	16740
GGCTAGCGGA CTTTTGAAGA TGGACTTTCTT GGGACTACGA AATTGACCT TTGTCAGAA	16800
GATGCAAGAG TTGCTTGCTG AAACAGAAGG TATTGATCTG AAAATTGAAG AAATCGATTT	16860
AGAAGACAAA GAAACGTTAG CTTTATTGCT CTCCTGGTAAT ACAAAAGGTA TCTTTCAATT	16920
TGAGCAACCA GGTGCCATTC GTCTGCTTAA GCGTGTGCAA CCAGTCTGTT TTGAAGATGT	16980
CGTCGCGACT ACTTCTCTAA ATCGACCGGG TGCTAGTGAC TATATCAATA ATTTTGTGGC	17040
AAGAAAGCAT GGGCAGGAAG AAGTGACTGT TCTGGATCCA GTACTGGAGG ATATTTGGC	17100

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TCCAACCTAC	GGCATAATGC	TCTATCAGGA	GCAGGTTATG	CAGGTTGCC	AGCGACTTGC	17160
CGGATTAGT	CTTGGAAAG	CCGATATTTT	GGTCGGCT	ATGGGGAAAA	AGGATGCCTC	17220
TGCCATGCAT	GAGATGAGGG	CTTCCTTTAT	TCAAGGTTCA	TTAGAAGCTG	GTCATACTGT	17280
GGAAAAGCA	GAGCAGGTCT	TTGATGTTAT	GGAGAAGTTT	GCAGGTTATG	GTTTAACAG	17340
GTCACACGCC	TATGCCTACT	CAGCCTTGGC	CTTCCAGTTG	GCTTATTTCA	AAACGCATTA	17400
TCCAGCCATT	TTTTATCAGG	TCATGTTAAA	TTCTTCCAAC	AGTGATTACT	TAATAGATGC	17460
ACTTGAAGCA	GGTTTGAAAG	TAGCCTCTCT	ATCCATCAAC	ACCATTCCCT	ATCACGATAA	17520
AATTGCCAAC	AAGGCATCT	ATCTAGTTT	GAAATCCATT	AAAGGAGTC	GTAATGATTT	17580
AGCTCTCTGG	ATTATTGAAA	ATAGACCTTA	TTCTAACATT	GAAGATTTA	TAGCTAAATT	17640
ACCTGAGAACAT	TATCTGAAAC	TTCCCTCTGCT	AGAACCTTG	GTTAAAGTTG	GTCTTTCGA	17700
TTCATTTGAA	AAAAATCGTC	AAAAAGTATT	TAATAACTTA	GCTAATCTAT	TTGAATTTGT	17760
GAAAGAGTTG	CGAAGTTTGT	TTGGAGATGC	TATTTATAGT	TGGCAGGAAT	CGGAAGATTG	17820
GACGGAACAA	AAAAAATTTT	ATATGGAACA	AGAGCTTTA	GGGATAGGTG	TCAGCAAACA	17880
TCCACTACAA	GCTATTGCAA	GTAAGGCTAT	TTACCCGATT	ACCCCAATCG	GAAATTTGTC	17940
AGAAAATAGC	TATGCTATTA	TCTTGGTTGA	AGTCAGAAA	ATAAAAGTGA	TTCGTACCAA	18000
AAAGGTTGAA	AATATGCCCT	TCTTACAGGC	AGATGATAGT	AAGAAAAAT	TGGATGTCAC	18060
TCTCTTTCA	GACTTATATC	GTCAGGTTGG	ACAGGAAATA	AAAGAGGGAG	CCTTCTACTA	18120
TGTAAAAGGA	AAAATACAAT	CACGTGATGG	CCGTCTCCAA	ATGATTGCAC	AAGAAATAAG	18180
AGAACCGAGT	GCTGAACGCT	TTTGGATACA	GGTAAAAAT	CATGAATCGG	ATCAAGAAAT	18240
TTCACGCATT	TTAGAACAAAT	TTAAAGGCC	AATCCCAGTC	ATCATCCGGT	ATGAAGAGGA	18300
ACAGAAAACC	ATCGTTTCTC	CCCATCATTT	TGTAGCTAA	TCCAATGAAT	TAGAGGAGAA	18360
ATTGAATGAA	ATCGTTATGA	AAACGATTTA	TCGCTAAAAA	TACGGAAAAT	AGAAGAATT	18420
TCAACGTTAA	TGTGGTATAA	TCAGTAAGAA	TGTTAAAAGA	AAAAGGAGCA	TAACCAATAT	18480
GAAACCTATT	GCTGTTTGA	CTAGTGGTGG	AGACGCCCT	GGTATGACG	CTGCCATCCG	18540
TGCAGTTGTT	CGTCAAGCAA	TTTCAGAAGG	AATGGAAGTT	TTGGTATCT	ATGACGGATA	18600
TGCTGGTATG	GTTGCCGGTG	AAATTCATCC	CCTAGATGCA	CCTTCAGTAG	GGGACATCAT	18660
TTCTCGTGGT	GGTACTTTCC	TTCACTCAGC	TCGTTACCCA	GAGTTCGCTC	AACTTGAAGG	18720
GCAACTTAAA	GGGATMTGAGC	AATTGAAAAA	ACACGGAATT	GAAGGTCTAG	TTGTTATCGG	18780
TGGTGACGGA	TCTTACCAACG	GGCCTATGCG	TTTGACTGAA	CATGGCTTCC	CAGCTATTGG	18840

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TCTTCCAGGT ACAATCGATA ACGATATCGT TGGTACTGAC TTTACAATCG GTTTTGACAC	18900
AGCGGTTACT ACTGCCATGG ACGCATATCGA TAAGATTCGT GATACATCAT CAAGTCACCG	18960
TCGTACTTTT GTAATCGAAG TTATGGGACG TAACGCTGGT GATATCGCTC TTTGGGCTGG	19020
TATTGCAACT GGTGCTGATG AAATCATCAT CCCTGAAGCA GGCTTCAGA TGGAAGATAT	19080
CGTAGCAAGC ATCAAAGCTG GTTATGAATG TGTTAAAAAA CACAATATTA TCGTCCTAGC	19140
TGAAGGTGTG ATGTCAGCGG CTGAATTGG TCAAAAACCTT AAAGAAGCTG GAGATACAAG	19200
CGACCTTCGT GTAACAGAAC TTGGACATAT TCAACGTGGT GGTTCTCAA CTGCGCGTGA	19260
CCGTGTTTG CGCTCACGTA TGGTGCACA TGCTGTTAAA CTTCTTAAAG AAGGTATCGG	19320
TGGTGTGCG GTTGGTATTC GTAACGAAAA AATGGTTGAA AATCCAATTC TTGGTACTGC	19380
AGAAGAAGGG GCATTGTTA GCCTTACTGC AGAAGGTAAG ATTGTGGTTA ACAACCCAGC	19440
TACAAA	19446

(2) INFORMATION FOR SEQ ID NO: 52:

- (i) SEQUENCE CHARACTERISTICS:
- (A) LENGTH: 16593 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 52:

TCGTAAATAT GCTCTGTTT TGGATTTGT TTCTTAATCT GTTTGGCAAG TGCCCTTCATC	60
ATAGAAATAG GACCACACAT ATAGACGGTT GCATGTTCGG GCACCTCTTT TTGTTCAAAA	120
TTAAGATAGC CGTCTTCGT ACTGTCGATT AGATGGAGTT CAAAATTAGG ATTTTTCTGA	180
GCATAGTTAC GGAGTAAATC TAGGTAGACT GCATTTCAT CTCCACGGAA GCTATAGTAG	240
AAGTGAACCT GTTTATCTAA AATAGGATGT TCACGGATGT AAGAGATGAA GGGGGTGATC	300
CCAATACCTC CAGCAATCCA AACCTGATTT TCTCGTCCTT CTTCTATGAT CATGTGTCCG	360
TAAGCTCTGT CTAGGGTTAC TTTGCTGCCG GCTTGAAGAT TATCATAGAT ATTCTTGGTA	420
TGGTCGCCCTG AAGTTTAAC AGTAAAGTAA AGAGTTTGAC CATGACCTCC TGAGATAGAA	480
AAGGGATGCG GAGCACTTC AAAGCCTCT TGAAAATCT TTAGAAAGGC AAATTGTCCCT	540
GATTGATAGT TGAAAGGTCT GCTAAGATGG ATTGAAATT CTCTAGTATC GTGATTTAAG	600
CGTTTGAGAT GGGTAATTTC CCCTAGATAG GGGAGGAAA TCTTTGATA TAGAAAAATG	660
ATATAAAAAC CAGCTAGTAA GCCTAAAAGG GCATAGCTAC CAACAAGAAA ACTTGAAGA	720
TTAAATGTAA GGAGACGATT GCCCATTATC ATGTAGATGT GAAAGAGTCC TAAAATATAG	780

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GCTAGGTAAA	CCAGGCGGTG	AATCCATCGC	CAAGCTTCGT	ATTGGATGTA	TTTGCCTAAA	840
TAGGCGACAA	GGATGATGCT	GGCAAAGATA	TAGATGGCAA	GATTGCCAA	CTGAGCAGCT	900
AAGCGAGAGC	CCCACAAACCC	GCCCCATACTA	AACTTATGAA	AGATTAGTAG	GATGATTGAG	960
AGAAAAGGCTG	TGAATTTGTG	GACGGGTGAG	ACCTTCTCCA	AACTGTGAA	CCAGCTTCT	1020
AGTAGTGGGA	GACGAGTGGC	TAGGATAAAA	GTCAGAGATA	GGCTTGTAA	AGCTAGTCCT	1080
GGAATCATGA	ATTGGGGAGA	AGTGTTCATC	CAAGTCAAA	GAGTCAGAT	AAAATAGCT	1140
ATGATAAAAGA	GTAGTCCTTT	GACTGATTTC	ATAGAAAATT	CCATTTCATT	TAGATTCGA	1200
TTTGTGTAA	ATAAATTTGT	TACATTTTAT	CATAGAAAAT	GTATGGTGTC	AAATTGAGGT	1260
CTATAAAATAT	CTACTCTCAT	CAAAAAACTC	TCCAATTGAA	CTGGAGAGTC	GCTGTTATA	1320
CTCAATGAAA	ATCAAAGAGC	AAACTAGGAA	GCTAGCCGCA	AGTTGCTCAA	AAACACTGTT	1380
TGAGGTTGCA	GATAGAGCTG	ACGTGGTTTG	AAAGAGATTTT	CGAAGAGTGT	TATTCTGCAG	1440
CTTGTGCCA	ACGTTGGCT	AGCATATGAG	ACAGGCTAGA	AATTGCTAGG	TTAAAGCTGA	1500
AGTAGATGAG	GGCAATCAGG	ATCTAAAGAC	TGAAGACCTG	CTCTGGTTCG	AAATAACGGC	1560
CCATGAGAAT	TTGGCTGGCT	CCAAAGAGTT	CTTGTAGGGC	GATAACAGAG	TAGAGGAGAC	1620
TGGTATCCTT	AATCACGGTA	ACAAACTGAG	AAATGATGCC	TGGTAGCATT	TTGCGGATGG	1680
CTTGTGGGAG	AATGATGAG	TAGAGGATTT	GGGCTGAGGT	GAAGCCCTGT	GACATTCCTG	1740
CTTCGTAATG	TCCCTTGCT	ACGGCATTGA	GACCGCCTCG	AATAATCTCA	GCCAAGGCTG	1800
CTGATGTTAA	GAGAGTAAAG	GCTGTAATAC	CTGCTGGTGT	GGATTTCATT	TTGAACACCA	1860
AAAAGATAGT	AAAATCCAG	AGAAGGTTGG	GAACGTTGCG	CACAAACTCG	ATATAAATAC	1920
TGGAAATAAT	GCGTAAGACA	GGATTTTGCG	CATTTCTCGT	GACAGCTAGC	ACCGTACCGA	1980
TGATAGTGA	GAGGATGATG	GCAATCACAG	AAATATAGAG	GGTCAAGCCA	AATCCTTTAA	2040
AGATAAAAGAC	TAGGTTATCT	GGGGTTAAAA	CTTCTAAAAT	AGATTCCATA	GTAACCTCCT	2100
AAAGTGAATA	GGCTTTTTG	TTGGCTTGCT	CCATCTGCG	ACCAAACTGG	GCAACAGGGA	2160
ACCATAGAGC	AAAGTAGAGA	AGAGCAGCAC	CTAAAAAGGC	TGGTATATAG	TTTCCGTTGA	2220
GAGCCGACCA	AGACTTAGTC	ACAAACATCA	AGTCTACTCC	AGAGATGATA	GCTACAGTAG	2280
AGGTGTTCTT	GATGAGGTTA	ACAATTTGGT	TGGTCAATGG	AGGGAGAATG	ATGCGGAAGG	2340
CCTGAGGCAA	GATAATCAAG	CGCATGGCAC	TGATATAGGT	AAAACCTTGC	GACAAGGCGG	2400
CCTCCATCTG	ACCACTAGGA	ATAGACTGAA	TCCCTGAACG	AATAACCTCA	GCGATATAAG	2460
CGCCCGTGATA	GAGTCCCACG	CAGAGAACGG	CTGTCCAATA	AATTGGAATC	ATGATGATAT	2520

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GGTCACTGAT AAGAGGTAGG CCATAAAAAA CAATAACAAA CTGCACCAAG AGGGGAGTAT	2580
TTTGGTAAAA TTCAACAAAG ATGCGAGCTA AAATGCGTAA AATTGGACGT TTACTGGTTG	2640
ACATGGCACC AAAGAAGATG CCCAAACCA TAGCGAGGAT AAAGGAACCA ACCGCTAGGG	2700
CAAGGGTGA GAGGAAACCA TTGAAAATT GTCCAAAATC CTGAAAATAG GCTGTCCAAG	2760
ATGATAAAATC TGTCAATGGGG TGTCCTCCTT AATCTGCAGT ATGGCTAGAT GGTTTGAGCT	2820
TGTAACGGTC ATAAAGTTTC TGCAAACATAC CATCCTTGCT CCATTTAGTA ACCAAGTTAT	2880
CAAGATAGTC GTTGAGCTCT GTATTTGATT TCTTGGTAAC AATACCGTAG TCAGATGGCT	2940
TGAAAATATC ATCTAGTAGT GCTGTCCGTT TACTAGTGTA GCCAGATAGA ATAGAGCGGT	3000
CAACGGAAAA GGTATCGATA CGATGAGCGT GCAGGGAAGT AATCAATTCT GGGTAGGAAC	3060
CAAGTTCGAC GAATTAAAC TTCAGACCTT TCTTTTACCC CAGTTCACTA ATCAGGCGTT	3120
GGGTGATAGA ACCTTGGCG ACTCCGATGG TTTTGCCCTT TAGTCCTCA ATCTTTTGAA	3180
TTTTGGCAGA TTTATTGACCA AAAATCCAG AAGCGTCTGT GTAGTAGGGAA CTGGTAAAGT	3240
TGTAGAGTTT TTTGCGTTG TCCGTGATGG TAAAGGTGCC GATATCCATA TCGACCTGTT	3300
CATTGTCATC AGCTACCATC TTGGCCAAGT CGGTTTCGAT ACCAGAATAA GTACCGGTCT	3360
TGGGATCTTT GTAACCAAAA TTGGGAACGT CTTGTTTGAC ACCGACAACC AGTTCGCCTC	3480
TTTTTGAAAT GTCTGCGATA CTTGTATCAG CCTGGACTGG TTTGGCAGCA GCAACCCCCA	3540
AAAGGCTAAT CAATAATGCT GATAAAAAGA ATTTTTTTTC ATAGGCGCCT CCTTATTGAA	3600
CTTTGTCACT TTCTGGTTG ATAATTTCG TGAGGAATTG TTGGGCACGA GGTTCGCTTG	3660
GATTGTCAAA AAAGTTATCG ACATCTGTG TATCTACTAA AACTTCTCCG TCGGCCATAA	3720
AGATAATGCG GTCCGCAACC TCTCGAGCAA AGCCCATTTC GTGGGTAACG ATGATCATGT	3780
TCATCCCATC ATGCGCCAGT TTCTGCATAA CTGCTAGAAC ATCTCCGATA GTCTCAGGAT	3840
CAAGAGCAGA TTCTGGTTCA TCAAAGAGGA GGAGTTCCGG ATGCATAGCA AGACCACGAG	3900
CGATGGCGAT CGCGTGTGTT TGTCACCCAG ATAGCATGGC GGGATAGGAA TCTTTCTTGT	3960
CCCACATATT TACAAATTCC AGATATTTTT GGGCGGTTTT TTCAAGCTTCT TTTTTATCAA	4020
TTCCCTAGAAC TTCAATGGGT GCAAGCGTTA CGTTTCTAA CACAGCTTGT TGTGGATAAA	4080
GGTTAAAATG TTGAAAAACCC ATCCCGACTT CCTTGCGAAG AGGTACCAA TCTTTCTGGC	4140
TGGCACCCAGC AACTTGGTGC CCATTGACTA GGAGACTTCC TTTGTCAACA GTCTCTAAC	4200
CATTGATCGT ACGGATAAGA GTGGACTTCC CAGAGCCAGA AGGTCCAAGC AGGACAACAA	4260
CTTGTCCCTT TTCAAAACGG AGATTGATGT TGCGGAATGC GTGGTAGTCT CCGTAATATT	4320

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TTTCGACGTT TTTAAATTCT ACTAAAGCCA TGAGAGATCT CTATTGTGTT ATATTTTATA	4380
ACACGGTTCT ACAATAAAAG AATGTTCTTG TCAAATCATA TCTGAAAAAA TTCACTATAG	4440
TGAAATAAGA ACAGGAAAAA TCGATCGGGA CAGTCAAATC GATTCTAAC AATATTTAG	4500
AAGTAGAGGT GTACTATTCT AGTTCAATA TACTATAAA TGTTATAAAA AAGCAATCTG	4560
GATAGAGAAA ACGTCTAAAT CATGTTATAA TGAAGCAATA GAATTCTTAG AAAGAGTGGA	4620
TGTCTTTTG ATAACACCTA CTTATGAATG GCAGTTGCC CTGCAGGTAG AAGATGCGGA	4680
TTTTACAAG ATAGCCAAGA AGGCTGGACT GGTCCTGAG GTGGCTCGGT TATTGTTGA	4740
GAGAGGGATT CAGAACCAAG AAAGTCTGAA GAAGTTTTA GAACCTTCCT TGAGGACTT	4800
ACATGATGCT TATCTGCTCC ATGATATGGA CAAGGCAGTG GAGCGGATTC GTCAGGCTAT	4860
TGAAGAAGGG GAAAATATTC TTGTTTATGG AGACTATGAT CGGGATGGCA TGACTTCGGC	4920
TTCTATTGTG AAGGAAAGTT TGGAAACAAT TGGTGCTGAG TGCCGAGTTT ACCTGCCAA	4980
TCGTTTACCC GATGGCTATG GCCCTAATGC TAGTGTATTA AAATACTTTA TCGAGCAAGA	5040
AGGGATTTC TTGATGTTGA CGGTGGACAA TGGGGTTGCT GGTCATGAGG CTATTGCATT	5100
GGCTCAGTCT ATGGGAGTAG ATGTCATTGT GACAGACCAT CATTCCATGC CTGAAACCT	5160
GCCAGATGCT TATGCTATTG TCCATCCTGA ACATCCAGAT CGGGATTATC CTTTTAAATA	5220
TTTGGCTGGT TGTGGAGTTG CTTTCAAGTT GCCTTGTGCC CTGTTAGAAG AAGTGCAAGT	5280
GGAAATTGCTT GATTGGTCG CTATTGGAAC TATTGCGAGAT ATGGTGAGTC TGACGGATGA	5340
AAATCGTATC TTAGTTCAAT ATGGCTGGGA AATGTTGGGT CATAACCCAGC GCATTGGCT	5400
GCAAGAAATG CTGGACATGG CTGGGATTGC TGCCAACGAA GTAACAGAAAG AAACGGTTGG	5460
TTTCCAGATT GCTCCTCGTT TGAATGCCCTT GGGTCGCTTG GATGATCCC ATCCTGCCAT	5520
TGATTTGTTG ACTGGATTTG ATGATGAGGA AGCGCATGAG ATTGCCCTTA TGATTCACCA	5580
GAAAAACGAA GAGCGCAAGG AAATCGTTCA GTCTATCTAT GAAGAAGCCA AGACCATGGT	5640
GGATCCTGAG AAGAAGGTC AGGTCTTGGC CAAGGAAGGC TGGAATCCTG GGGTTCTAGG	5700
AATCGTGGCT GGTCGTTAT TGGAAAGAATT GGGACAGACA GTCATTGTT TTAATATAGA	5760
AGACGGTCGT GCCAAGGGCA GTGCTCGTAG TGTGGAAAGCG GTCGATATTT TTGAAGCTCT	5820
GGATCCCCAT CGAGACCTCT TCATGCCCTT TGGAGGTCAT GCAGGTGCAG CGGGTATGAC	5880
GCTGGAAGTT GAGCAACTCT CAGATTATC TCAGGTTTG GAAGATTATG TTCGTAAAAA	5940
AGGTGCAGAT GCTGGTGGCA AGAATAAGTT AACCTAGAT GAAGAGTTGG ATTTGGAGGC	6000
ACTTAGCTG GAAACGGTCA AAAGTTTGAGCT CCTTTTGAA TGGATAATCA	6060

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GAAACCTATT	TTTTATATCA	AGAATTTCA	GGTCGAAAGT	GCTCGTACTA	TGGGGCAGG	6120
TAATGCCCAT	CTAAAGCTGA	AAATTCCAA	GGGTGAGGCG	AGTTTGAAAG	TGGTAGCCTT	6180
TGGTCAAGGC	AGATGGGCGA	CAGAGTTTC	TCAAACCAAG	AATCTAGAGT	TAGCGGTTAA	6240
ATTGTCTGTC	AACCAATGGA	ATGCCAAC	TGCCCTCCAG	TTGATGATGG	TGGATGCGCG	6300
AGTGGAAAGGT	GTTCAACTTT	TTAACATTG	TGGAAAAAT	GCAGTCTTGC	CAGAAGGTGT	6360
TCCAGCTTG	GATTTCCCTG	GAGAACTGCC	AAATCTTGC	GCTAGTGAAG	CTGTTGTCGT	6420
AAAAAACATT	CCAGAGGATA	TTACTCAGCT	GAAGACCATT	TTTCAGGAAC	AGCATTCTC	6480
TGCTGCTAT	TTCAAAAATG	ATATTGACAA	GGCTTATTAT	CTGACAGGTT	ATGGGACTAG	6540
AGATCAGTTT	GCCAAATTGT	ACAAGACTAT	TTACCAAGTTC	CCAGAGTTG	ATATTGCTA	6600
CAAGCTGAA	GATTTGGCTG	CATATCTAA	TATTCAACAA	ATCTTGCTGG	TCAAGATGAT	6660
TCAAGTATT	GAAGAACTAG	GCTTTGTGAC	GATAAAAGAT	GGTGTGATGA	CAGTCATAA	6720
AGAGGGCCA	AAGCGGGAGA	TAGGAGAAAG	TCAAATTTAC	CAAATCTCA	AACAAACCGT	6780
TAAAGACCAA	GAAATGATGG	CGCTGGGTAC	GGTGCAAGAA	ATTTATGATT	TTTGATGGA	6840
AAAAGAGTAG	AAGTTAGGAA	AGAGTTGGGA	AATCAACTCT	TTTTGAAAAA	CAGACCTTC	6900
TTTGAAAAAT	CATCAAAAAA	ATGGTATAAT	GGTAGGAAAA	GATTCGGCTG	AAAGTATCAG	6960
AACTTTAGA	ATAAGAGGGT	AGAATTGCC	TATAATCAAG	ATAAACTAAG	ATTTGGAGG	7020
AAAAATGAGT	AATATCAGTT	TAACAACACT	TGGTGGTGTG	CGTGAGAATG	AAAAAAATAT	7080
GTACATTGCT	GAAATTGGAG	AGTCCATT	TGTTTTGAAT	GTAGGGTTAA	AAATATCCTGA	7140
AAATGAACAA	TTAGGGTCG	ATGTGGTGAT	TCCAAACATG	GATTACCTTT	TTGAAAATAG	7200
CGACCGTATT	GCTGGGTTT	TCTTGACCA	CGGGCATGCG	GATGCCATTG	GTGCTCTACC	7260
GTATCTCTG	GCAGAGGCTA	AAGTTCTGT	ATTTGGGTCT	GAGTTGACCA	TTGAGTTGGC	7320
AAAGCTCTT	GTCAAAGGAA	ATGATGCCGT	TAAGAAATT	AATGATTTCC	ATGTCATTGA	7380
TGAGAAATCG	GAGATTGATT	TTGGTGGGAC	AGTGGTTTCC	TTCTTCCCTA	CGACTTACTC	7440
CGTTCCAGAG	AGTCTGGGAA	TTGTCTGAA	GACATCGGAA	GGAAGCATCG	TTTATACAGG	7500
TGACTTCAAA	TTTGACCAA	CGGCTAGTGA	ATCTTATGCA	ACTGATTTG	CTCGTTGGC	7560
AGAGATTGGT	CGTGACGGCG	TCCTGGCTCT	CCTCAGTGT	TCGGCCAATG	CAGACAGCAA	7620
TATTCAAGTG	GCTAGTGAA	GTGAAGTTAG	GGATGAAATT	ACCCAAACTA	TTGCTGACTG	7680
GGAAGGTCGT	ATCATCGTTG	CAGCTGTTTC	CAGTAATCTT	TCTCGTATTG	AGCAGATTTT	7740
TGACGCTGCG	GATAAAACAG	GTCGACGTAT	CGTCTTGACA	GGATTTGATA	TTGAAAATAT	7800
CGTCCGCACA	GCGATTGTC	TTAAGAAGTT	GTCTTTAGCC	AACGAAATTC	TTTGATTAA	7860

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GCCTAAAGAT ATGTCTCGCT TTGAAGACCA TGAGTTGATT ATTCTTGAGA CAGGTCGTAT	7920
GGGTGAGCCT ATCAATGGAC TTCGTAAGAT GTCGATTGGT CGCCATCGTT ATGTAGAAAT	7980
CAAGGATGGG GACCTAGTCT ATATTGCTAC GGCTCCGTCT ATTGCTAAAG AAGCCTTGT	8040
TGCGCGTGTG GAAAATGTA TTTATCAGGC AGGTGGGGTT GTCAAATTGA TTACCCAAAG	8100
TTTACATGTA TCAGGGCACG GAAATGTGCG TGATTTGAG CTGATGATCA ATCTTTGCA	8160
ACCTAAGTAC CTCTCCCTG TCCAAGGGGA GTATCGTGAG TTGGATGCTC ACGCTAAGGC	8220
TGCCATGGCA GTTGGGATGT TGCCAGAACG CATCTTCATT CCTAAAAAGG GGACGACCAT	8280
GGCTTACGAG AATGGAGACT TTGTTCCAGC TGGATCGGTT TCAGCAGGAG ATATCTTGAT	8340
TGATGGAAT GCCATTGGTG ATGTTGGAAA TGTTGTTCTT CGTGACCGTA AGGTCTTGT	8400
AGAGGATGGA ATTTTCATCG TGGCTATTAC AGTCAACCGT CGTGAGAAGA AAATTGTGGC	8460
TAGGGCTCGT GTTCACACGC GTGGATTTGT TTATCTCAAG AAGAGTCGG ATATTCTCCG	8520
TGAAAGTTCA GAATTGATTA ACCAAACGGT AGAAGAGTAT CTTCAAGGAG ATGACTTTGA	8580
CTGGCAGAT CTCAAAGGTA AGGTTCGTGA CAATCTGACC AAGTACCTCT TTGATCAAAC	8640
CAAGCGTCGC CCAGGCCATT TACCAAGTAGT CATGGAAAGCA AAAATAATCGT TGAAATAAAC	8700
AGAGAGAAAG TCGAGTTTCG GCTTTTCTT ATAGAAAAT AGAAGGAGAA AATCATGGCA	8760
GTGATGAAA TCGAGTATTA CTCACAAGTA TTGGATATGG AGTGGGGGT GAATGTCCTC	8820
TACCCCTGATG CCAATCGAGT GGAAGAACCA GAGTGTGAAG ATATTCCCGT CTTGTACCTT	8880
TTGCACGGGA TGTCTGGAAA TCATAATAGT TGGCTTAAGC GGACCAATGT AGAACGCTTG	8940
CTTCGAGGAA CTAATCTCAT CGTTGTTATG CCAATACCA GCAATGGTTG GTACACCGAT	9000
ACCCAGTATG GTTTGACTA CTACACGGCT CTAGCAGAGG AATTGCCACA GGTTCTGAAA	9060
CGCTTCTTC CTAATATGAC GAGCAAGCGT GAAAAGACCT TTATCGCTGG TCTTTCTATG	9120
GGAGGCTACG GCTGCTTCAA ACTGGCTCTT ACACAAATC GTTTTCTCA TGCAAGCTAGT	9180
TTTCAGGTG CCCTCAGCTT TCAAAACTTT TCTCCTGAAA GTCAAAATCT GGGAAAGTCCA	9240
GCCTACTGGA GAGGTGTTTT TGGAGAGATT AGAGACTGGA CAACTAGTCC CTATTCTCTT	9300
GAAAGTCTGG CTAAAAAATC GGATAAAAAG ACCAAACTTT GGGCGTGGTG TGGCGAACAG	9360
GATTTCTTGT ACGAAGCCAA TAATCTCGCA GTGAAAATC TCAAAAACT AGGTTTTGAT	9420
GTGACCTATA GCCATAGCGC TGGAACCTCAC GAGTGGTACT ACTGGGAAAA ACAATTGGAA	9480
GTTTTTTAA CAACCCCTACC AATTGATTTC AAATTAGAAG AGAGACTGAC TTAGTTGAA	9540
CTTCAGCATA GGGGGAGTAG AACTAAAATA AAATATGTTT TCACTAGACT TTTCAAACGm	9600

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AAGTAGTAGA ATAGTAATAA AATACTGGAG GAAAGAGAGT AGGAAATGTA CCGTTATCAA	9660
ATTGGCATTG CCACATTAGA ATATGATCAG TTTGTCAAAG AACATGAATT AGCCAATGTA	9720
TTACAAAAGTA GTGCTTGGA GGAAGTTAAC TCTAATTGGC AACATGAGAA GTTTGGTGT	9780
TACAGGGAAG AAAAATTAAGT GGCGACAGCT AGTATTTGA TTAGAACTCT TCCGCTAGGC	9840
TATAAAATGT TTTACATCCC AAGAGGACCT ATATGGATT ATGGGATAA AGAACTCTTG	9900
AATTTGCCA TTCAGTCTAT TAAGTCCTAT GCTCGCAGTA AGAGAGCGGT TTTTGTGACT	9960
TTTGACCCAA GTATTGCCT ATCTCAAAGT TTAATCAATC AGGAAAAGAC AGAATTCCT	10020
GAAAATCTGG CTATTATTGA TAGTTGCAA CAAATGGAG TAAGGTGGTC AGGAAAAACG	10080
GAGGAAATGG GAGACACCCT TCAACCTCGT ATTCAAGCGA AAATATACAA GGAAAATTT	10140
GAAGAAGATA AACTTCCAA GTCAACAAAA CAGGCTATTG GAACAGCAG AAACAAAGGG	10200
CTTGAGATTC AATATGGTGG ACTGGAACTA TTAGATTCA TTTGGAGTT GATGAAAAAA	10260
ACTGAGAAGC GAAAAGAGAT TCATTGAGG AATGAAGCCT ATTATAAAA ATTGTTAGAT	10320
AATTTTAAGG ACAAGGCCTA TATCACCTTG GCCACCTTGG ATGTTCTAA ACGTTCGCAA	10380
GAGTTAGAAG AACAGTTAGC GAAAATAGA GCCTTGGAG AGACCTTAC TGAGTCGACT	10440
CGAAACTCAA AAGTAGAAGC GCAGAAGAAG GAAAAGAAC GTTTGTTAGA GGAATTGACC	10500
TTCTTGCAGG AATATATAGA TGTAGGTCAA GCGAGAGTTC CTTTAGCGGC TACTTTGAGT	10560
TTGGAATTG GTACTACCTC TGTCAATATA TATGCTGGTA TGGATGATGA TTTTAAACGT	10620
TACAATGCAC CAATTTAAC TTGGTATGAA ACGGCTCGCT ATGCCTTGA ACGAGGTATG	10680
ATCTGGCAA ATTTAGGTGG TGTGAAAAC TCTCTCAATG GTGGACTTTA TCATTTAAAG	10740
GAAAATTTA ATCCAACGAT TGAAGAATAC TTGGGTGAAT TTACAATGCC CACTCATCCT	10800
CTCTATCCTC TGTTAAGACT TGCTCTTGAT TTCCGTAAAA CATTAAGAAA AAAACATAGA	10860
AAGTAAGTAT ATGGCACTAA CAACACTCAC GAAAGAAGAG TTTCAGACTT ATTCTGATCA	10920
GGTTTCTTCT CGTTCTTTA TGCAATCTGT CCAGATGGGG GATTTGCTAG AAAAAAGAGG	10980
GGCTCGAAATT GTTATCTTG CTTTGAAACACA AGAAGGAGAA ATTCAAGTTG CAGCTCTGGT	11040
TTATAGCCTG CCCATGCTGG GTGGCTGCA TATGGAACCTC AATTGGGGC CGATTATAC	11100
CCAACAAGAT GCTCTCCAG TTTTTATGC AGAGTTAAA GAATATGCC AGCAAAATGG	11160
TGTATTAGAG TTGCTTGTAA AACCTATGA AACTTATCAA ACTTTTGATA GCCAAGGTAA	11220
TCCAATAGAT GCTGAGAAAA AAAGTATTAT TCAAGATTG ACTGATTGTTAG GTTATCAATT	11280
TGATGGCTTA ACAACAGGTT ACCCAGGTGG AGAACCCAGAT TGGTTATACT ATAAAGATTT	11340
AACTGAATTAA ACTGAAAAGA GTTTGCTTAA AAGTTTTAGC AAAAAGGGTA AACCCCTGGT	11400

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GAAAAAGGCT GAAACCTTG GCATTCGGTT GAAAAAGTTA AAACGTGAAG AACTATCGAT	11460
TTTTAAGAAT ATAACAAAAG AAACCTCTGA ACCTAGAGAA TATAGTGATA AAAGTTAGA	11520
ATATTATGAG CATTTCATG ATACTTTGG AGAACAGCG GAGTTCTCA TAGCAAGCTT	11580
AAATTTTCG GACTATATGA GCAAATTGCA AGGTGAACAA AGTAAACTAG AAGAAAACCTT	11640
GGACAAGTTG CGACTTGATT TGAGTAAAAA TCCTCATTCT GAGAAAAAAC AAAATCAACT	11700
GAGAGAATAT TCTAGTCAT TTGAAACGTT TGAAGTTCGA AAAGCAGAAG CGCGAGACTT	11760
GATTGAAAAA TATGGAGAAG AAGATATTGT TTTAGCTGGG AGTTTATTTG TTTATATGCC	11820
TCAGGAAACG ACTTATCTCT TTAGTGGTTC CTACACTGAG TTTAATAAGT TCTATGCC	11880
TGCAC TGCTT CAAAAATATG TTATGTTGGA AAGCATAAAA CGTGGAAATAC CTAAATACAA	11940
CTTCCTAGGC ATTCAAGGGA TTTTGATGG AAGTGTGGT GTTTTGCCTT TTAAACAGAA	12000
TTTTAATGGC TATATTGTAC GCAAAGCAGG TACCTTCGTT TACCATCCAT CGCCTTTAAA	12060
ATACAAAGCT ATCCAGTTAC TCAAAAAAAT AGTAGGACGT TAAGATGAAA AAGTCAGTAT	12120
TTAGATTTCT TTTAGCTTCT TTTAGTAAAAA TAATTCTTAT TTGCTAGAAA GGTGGAGAGA	12180
CATGCGCTGG CTTTTTCGTT TGATAGGGC TTTCTTTCT TTTGTGTGGC GTTTGTTTG	12240
GCGCTGGTT TGGATAGTTG TGCTCTTATG TGTGCTTGCT TTGGACTTC TCTGGTATCT	12300
GAACGGAGAT TTTCAAGGAG CGCTAAAGCA AGCAGAACGG TCAGTAAAAA TTGGTCAACA	12360
AAGTATTGAC CAATGGGAGA AAACAGGGCA ACTGCCTAAG TTAAGCCAGA CAGATAGTCA	12420
CCAGCATTCT GAAGGAAGGT GGGCACAGGC CTCTGCTCGT ATTTACCTGG ATCCGCAGAT	12480
GGATTCACGC TTTCAAGGAG CTTTATTAGA AGCAATCCAG AACTGGAATC AAACGGTGC	12540
TTTTAACTTT GAACTCGTGA CTGAGTCTAG TAAGGCGGAT ATTACGGCTA CGGAGATGAA	12600
CGACGGAGGC ACTCTGTGG CAGGAGAGGC GGAAAGTCAA ACTAATCTCT TAACAGGGCA	12660
ATTCTTGTCG GTAACGGTGC GGTTGAATCA TTATTATTTG TCCAATCCAT ACTATGGCTA	12720
CTCCATATGAA CGCCTGTCC ATACGGCAGA ACATGAGTTA GGTGATGCCA TTGGCTTGGA	12780
CCATACAGAT GAGAAGTCTG TCATGCAACC AGCAGGTTCC TTTTATGGTA TCCAGGAAGA	12840
GGATGTTGCA AACCTCCGAA AAATATATGA GACTAGTGGAG TAGGGTACTA TCTTTCCCTA	12900
CTTTTTTGCA TATAATGGAA CTATGAACAA CTTGATTAAA TCAAAACTAG AGCTCTTGCC	12960
GACCAGCCCT GGTTGCTACA TTCATAAGGA TAAAATGGC ACCATTATCT ATGTAGGAAA	13020
GGCTAAAAAT CTGCGTAATC GAGTACGGTC CTATTTCTGTT GGAAGTCATG ATACCAAGAC	13080
AGAGGGCTCTG GTGTCTGAAA TTGTGGATTT TGAATTATTGTTACGGAGT CTAATATTGA	13140

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GGCACTTCTC	CTAGAAATCA	ACCTGATCAA	GGAAAACAAG	CCCAAGTACA	ATATCATGCT	13200
CAAGGATGAC	AAGTCTATC	CTTTCATCAA	AATCACCAAT	GAGCGCTATC	CACGCTTGAT	13260
TATCACTCGT	CAGGTCAAAA	AGGACGGAGG	TCTTTATTTT	GGACCCATTC	CCGATGTGGG	13320
GGCAGCCAAT	GAAATCAAGC	GGTTGCTGGA	TCGGATATTC	CCTTTTCGTA	AGTGTACCAA	13380
CCCGCCCTCT	AAGGTCTGTT	TTTATTACCA	TATCGGCCAG	TGTATGGCCC	ACACCATCTG	13440
TAAGAAGGAT	GAGGCTTATT	TCAAGTCTAT	GGCCCAGGAG	GTTGCTGATT	TTCTGAAAGG	13500
TCAGGATGAC	AAAATCATCG	ATGATCTCAA	GAGTAAAATG	GCAGTAGCAG	CACAAAGTAT	13560
GGAGTTTGA	CGTGGCGCGG	AATACCGTGA	CCTGATTCAG	GCTATTGGAA	CGCTTCGAAC	13620
CAAGCAACGG	GTCATGGCGA	AAGATTTGCA	AAATCGCGAT	GTCTTGCGCT	ACTATGTGGA	13680
TAAGGGCTGG	ATGTGTTGTC	AGGTTTTCTT	TGTCGTCAG	GtAAGCTCAT	CGACCGCGAT	13740
GTCAATCTCT	TCCCCACTT	CAATGATCAA	GATGAGGATT	TTTTGACCTA	TGTAGGACAA	13800
TTCTATCAAG	AAAAATCTCA	TCTAGTTCCC	AATGAGGTAC	TGATTCCGCA	GATATTGACC	13860
AAGAAGCTGT	CAAGGCTTG	GTGGATTCCA	AGATTCTTAA	GCCTCAACGT	GGACAGAAAA	13920
AACAACCTGGT	CAATCTAGCC	ATAAAAAAATG	CTCGTGTAG	TCTAGAGCAG	AAGTTCAATC	13980
TGCTAGAAAA	ATCTGTCGAA	AAGACTCAAG	GACCTATTGA	AAATCTAGGG	CGTTGCTCC	14040
AAATCCCGAC	CCCAGTACGT	ATCGAGTCCT	TCGATAACTC	TAATATCATG	GGAACTAGCC	14100
CTGTTTCGGC	TATGGTGGTC	TTTGTCAACG	GTAAACCGAG	TAAGAAGGAT	TACCGTAAGT	14160
ACAAGATAAA	AACGGTTGTT	GGACCAAGACG	ACTATGCCAG	CATGAGAGAG	GTCATTGCA	14220
GACGCTATGG	TCGAGTACAG	CGTGAGGCTT	TGACTCCTCC	AGATTTGATT	GTGATTGATG	14280
GGGGGCAAGG	TCAAGTCAT	ATCGCTAAGC	AGGTTATCCA	AGAGGAACGT	GGCTTGATA	14340
TTCCAATTGC	TGGGCTGCAA	AAGAATGATA	AGCACCAAC	CCATGAATTG	CTCTTTGGAG	14400
ATCCGCTTGA	GGTGGTGGAT	TTGTCTCGCA	ATMCTCAGGA	ATTTTCCTC	CTCCAACGCA	14460
TCCAAGATGA	GGTGCACCGC	TTTGCTATCA	CTTTCCACCG	CCAACGCGC	TCCAAAATT	14520
CTTTCTCATC	TCAATTGGAT	GGGATTGACG	GTCTGGGACC	TAAACGCAAG	CAGAACATCTTA	14580
TGAAGCATT	CAAGTCTTG	ACCAAAATCA	AGGAAGCCAG	TGTGGATGAG	ATTGTCGAAG	14640
TTGGGGTACC	TAGAGTCGTT	GCAGAGGCTG	TGCAAAGAAA	GTTGAACCCG	CAGGGAGAAG	14700
CCTTGCTCTA	AGTAGCAGAA	GAAAGAGTAG	ATTACCAAAC	GGAAGGAAAC	CACAATGAAC	14760
CATAAAATCG	CAATTTTATC	AGATGTTCAT	GGCAATGCGA	CGCGCCTAGA	AGCAGTGATT	14820
GCAGATGCTA	AAAATCAAGG	GGCCAGTGAA	TATTGGCTTC	TGGGAGATAT	TTTTCTTCCT	14880
GGTCCAGGCG	CAAATGACTT	AGTCGCCCTG	CTAAAGGACC	TTCCTATCAC	AGCAAGTGTT	14940

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CGAGGCAATT GGGATGATCG TGTCCTTGAG GCTTTAGATG GGCAATATGG CTTAGAAGAC	15000
CCACAGGAAG TTCAGCTCTT GCGTATGACA CAGTATTGAGA TGGAGCGAAT GGATCCTGCA	15060
ACGATTGTCT GGCTACGAAG CTTGCCTTG CTGGAAAAGA AAGAAATTGA CGGATTGCC	15120
TTTTCTATCT CTCATAATTT ACCTGACAAA AACTATGGTG GTGACTTGCT AGTTGAGAAT	15180
GATACAGAGA AATTTGACCA ACTGCTAGAT GCGGAAACGG ACGTGGCAGT TTATGGTCAT	15240
GTTCACAAGC AGTTGCTTCG TTATGGAAGT CAAGGGCAAC AAATCATCAA TCCAGGGTCC	15300
ATTGGCATGC CCTATTTAA TTGGGAGGCG TTAAAAAATC ACCGTTCCCA GTATGCCGTG	15360
ATAGAAGTTG AAGATGGGGA ATTACTCAAT ATCCAATTTC GTAAAGTTGC TTATGATTAC	15420
GAAGCTGAGT TAGAATTGGC CAAGTCCAAG GGGCTTCCCT TTATCGAAAT GTATGAAGAA	15480
CTGCGTCGTG ACGATAACTA TCAGGGGCAC AATCTGGAAT TATTAGCCAG CTTAATAGAA	15540
AAGCATGGGT ATGTAGAGGA TGTGAAGAAT TTTTTGATT TTTTGTAAGA GTTTCTAAA	15600
ATAGCCAATG CAAACTAAAA AAGCGATTTG CTGGTCCAAT CGCTTTAGT ATATCTTATA	15660
CTCAATGAAA ATCAAAGACC AAACTAGGAA GCTAGCCGTA GGTTGCTCAA AGCACAGCTT	15720
TGAGGTTGCA GATAAAGCTG ACGTGGTTTG AAGAGATTT CGAAGAGTGT TATTGTAACT	15780
GAGATTGATC TGGGAGGTAA GAACCACCTA GATAGGTATT GCTGAGTTT TCAAGGGTTC	15840
CGTCTTGATA GAGTTCTTG AGCGCTTAT CAAATTGCTC TTTAAACTCT TTTGGTCGC	15900
TTGAGAAAAT GATATAATTG CTGGGCTAT CTGCAGAAGG TAAATCAACG ACTGAGAGGT	15960
CTAAACCACG GTCCTTGATA ATCTTTGAA CGGATACCTT GTCAAAACT AGGAAATCAA	16020
ACTCTCCGTT AGCAAGGTCT AGGATTGTT TACCAATATC CTCACCAGAA AAATTAATTG	16080
TAGCGGGATT ATCAGTGTGT TTCTGATTCC AGTTATTGAT GAATTGAGCG TTAGAAGTTC	16140
CGGTATCCTC TTGTGTTGTT TTACCAAGCGA TCTGGTCAG AGAAGTCAAA GGATTTTCT	16200
TGTTGCTGAC AAGGACGAGG GGATTGTTGG AAATTGGAAG CGAGTAAAGG TATTTTCAG	16260
CACGCTCTTT TGTGTAACTC AAGTTATTGG CCGCAGCCTG ATAGTGACCA GAATCAAGTC	16320
CTGGGAAGAT GCTCTCCCAG GCGGTTCTTT GGAATTGAAT CTCGTAGTCG CTGAGTTTT	16380
CATCTACTGC CTTAAAAACT TCGATATCAA AGCCTGTCAG ATTGCCCTG TCTTCGTAGT	16440
CAAATGGTGG CACGTCGCCA GCTGTAGCAA GGACGATTGT CTTTGAGCG CTAGTCTCTT	16500
TGGGTGAGC TTGATTCTCA CAGGCAACCA AAAATGGTAG GATAGCTAGT AATAGGCTAA	16560
ATTTTTTCAT ACTGCTCCA TTCAATGTA AAG	16593

(2) INFORMATION FOR SEQ ID NO: 53:

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(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 3510 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 53:

GGGATATCCT TATATCCTTG TTCCCTGGAAC CATTGTGGGA ATTGCTCAAC AGTTTTTCA	60
CCTTGAATTC CTGGTGCAAT GACAGTAAGA ATTTCGAAAT CACGATCTGG TTTCGCCGCT	120
AGTTCATCA ACTCTGGCAT ACTTTCTTG CATGGACCAC ACCATGAAGC CAAAAACTTC	180
AAGTAAACCT TTTTACCCCTT AAAATCAGAT AACTTAACCTT CTTTGCCATC CATGGATTGC	240
AATGTGAAGT CTGGAGCATC TTTTCCAACA GCAATTGTT GTACAGTCGT TTGTTGTTT	300
GGCTGTTG TGCTGTTGAGT CTTTTTAGTT TCTTCCTCAC CACAGGCCAT CAATACAAC	360
AATGACAAGA GACTTAAGCC AGCAAACATT ACTTTTTCA TTTGTCCTCC TTTTATTCAA	420
AATTCCAGCT AGAACATTAA CTTGTCCTAA TAGTAACAAA ATTCCCATTAA AAACAATGAG	480
GAAACCACCA ATTTCTTTA TAGCATCAT ATGACGCTTG ATTTTACTAA AATATGGCAT	540
GAATGACCT GAAGCTAGTG CCAATACCAA GAAAGGAAGG GCCATGCCAG AGTGTAAATG	600
AGAGTATAAA TCGCTCCTTG CCAAGCGCCA TTGCTCCAG AAGCCGCAAG TGCTAAAACA	660
GAACCTAAAA CTGGACCAAT ACAAGGTGTC CAACCAAAGC TAAAGGTAAT ACCAACTAAA	720
AAAGCTGACC ATAACGATT AGAATCTGAT TTTTAAAGG TAAACTTTT TTGAACTTCT	780
AATTTCTTC AATGAAAAAT TTCCATCTGG TGAAGACCCA AAATGATAAT AATAGCTCCC	840
ATGCCATATC, GAAACCAATT TGCATAGAGA ATATGACCAA AGTAACCAGC ACCAAAGCCT	900
AGAATAAAGA AAATGAGAGA GATACCAGCG ATAAAGCAA GTGTTGCAAT CAAGCCTGAC	960
CAGAGAACCT TTCTCCAAA CAAAGAAAAG CTTTTGCAC TTTCTGATC ATCCAATAAA	1020
ATCCCAGCAT AGACTGGCAG AAGAGGAAAA ATACAAGGAG AAAAAAGGA TAAACACCT	1080
GCTAGAAAAA CAGAGATTAA AAATACTATC GTTCTCAATA AAGAACCAAC TTTCTTAATA	1140
ATTCTAATCC TATTTTACTA TATTCAATT TATTTGTAAG CTTTCTGCTA CGAAAATCG	1200
TATCGGGCAC TATTGGACCA ATCTTTCTT TTGCTAGTCA AGCGGGATCT TATCCCCCAA	1260
AATAGCCAA AAGCAACGAC AAGGATTACT CATCGCTGCT TTTGTGAACG AAAATGTCTT	1320
TTAGGTCTGA CATTTCATAA ATCATGTTT ACTTGAGTTT GTCAAGGATT GCTTTAAGCT	1380
CCTCTACTAG TTTAGTTCT GTCTCTGCTG AGCCATTTTC TTCTTTACG AAATCAAGGG	1440
TTTCTGGAG AAGGTTTGG GCTTGGCAA GGACTTTTTT ATCCGTTTT TCTGCATCTA	1500

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GCTGTCCTAG AACCTTGATC AATTCCGTGC TTAATTGCTG GATTTCTGAC TCTTTCTTAC	1560
GGCGAATCAG CCAGAAGGCA ATCACGCCA GGAGGGCAAG TAGACTGACC ACAATCACTC	1620
CTGCGGAAAC TGAGTTTGTT TCAGTCATCT TATCTGAATC CTTACTATCT TCCGTTCTT	1680
GTTCCTGATC CTTCTTGTC TGTCAGGCT TGCTGTCGCT AGCATTGCT TTCAACATCTT	1740
TGAGAGAGTC CAAGGCAGCC CACCCCTTCAC AGACTCTACT GCAGTATGCA GACCTTACTC	1800
TGTCAAGGCA CTATCTTCGG GAGCTTTTG AGCATCTAGG AGGACAGCCT TGGTTGCATC	1860
GATTTTCGGA TCAGATACTG TTGCCAAAGC TTTCAAGCGT TGGTCTAACT CTTGACTCAA	1920
GGCACCGAGT TCAGACTTGT CAACCTGTC TTGAGCTTGT GTGCTCGTT AGCTAGCCGA	1980
AGCGCTTGCT ACCACTCTAG GATCTTGAGT CGGAGCTGAG CTTGGAGCTG GGACAGGGCT	2040
TGCAGGTTGA CTAGGAACAG TTATGGTATA TTGAAACTAG AATAGTACAT ATGGACTTCT	2100
AAAACATTGT TAGAATTGCA TTTTACTGTC CTGATCGATT TGTCTTATTCTTATTC	2160
TTACTATAAT AACCGATGGT GTGGTTAATG TTGGTAAGAG AAACCTCTGA AACCAAGCTT	2220
CAAAAAAGTC GCTCGTCATC GTCTCTTCGT AAGTCATGG AGCGATTAAT TCACCATTTG	2280
TTAGACCTGC AACCAAAGAA ATCCTCTGAT ATCTTCTTCC AGATACTTTG CCTCTTATTA	2340
ACTGACCTTT TAATGAGCGA CCATATTCTC GATAAAAATA AGTATCGAAT CCTGTTCGT	2400
CAATCTAAC ACAGTGCTAGG TGCTTTAAC TATTTAAATT CTTAAGAAAT AAGGCTACTT	2460
TTTCTGGTC TTGTTCATAG TAGGTGTGGT TCTTTTTTTC GAGTGTAGCC CATAGCTTTG	2520
AGCGCATAGT GGATGGTAGT TGGATGACAG CCAAACCTAG AAGCTATTTC AGTCAAATAA	2580
CCrTCTGGAT TGTCAGTAAG ATAGTTTTTA AGTCTATCTC TATCAACTTT TCTTGGTTTT	2640
GTTCCTTTTA CTTGGTGGTT TAGCTCTCCT GTTTCTCTT TTAGCTTTAA CCAGCCATAA	2700
ATGGTATTAC GTGAGATTTG GAAAACGTGT GATGCTTCTG TTATACCTACC TATTCGCTCA	2760
CAATAAGAGA GAACTTTTT ACCAAAATCT ATTGAATATG CCATAAGAAG ATTATACCA	2820
ATTGTGTACT ATTTTGGTT CATTCACTA TAACACAAAA TAGATTATTA TTACATAACA	2880
AAAAAGAGGT CTAAACCTCT TAATCTCAATT ACTCCGCCAG TAGGACTCGA ACCTACGACA	2940
TCATGATTAA CAGTCATGCCG CTACTACCAA CTGAGCTATG CGCGATTAAA GCTAAGCGAC	3000
TTCCCTATCT CACAGGGGGC AACCCCCAAC TACTTCCGGC GTTCTAGGGC TTAACCTCTG	3060
TGTTGGCAT GGGTACAGGT GTATCTCCTA GGCTATGTC ACTTAACCTCT GAGTAATACC	3120
TACTCAAAAT TGAATATCTA TTCAATTAA GAAAACCGTT CGCTTTCATA TTCTCAGTTA	3180
CTTTGGATAA GTCTCGAGC TATTAGTATT AGTCCGCTAC ATGTGTCGCC ACACCTCCAC	3240

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TTCTAACCTA TCTACCTGAT CATCTCTAG GGCTCTTACT GATATATAAT CATGGAAAT	3300
CTCATCTTGA GGTGGkTtCA CACTTAGATG CTTCAGCGT TTATCCCTTC CCTACATAGC	3360
TACCCACCGA TGCCCTTGGC AAGACAACATG GTACACCAGC GGTAAAGTCCA CTCTGGTCT	3420
CTCGTACTAG GAGCAGATCC TCTCAAATTT CCTACGCCCG CGACGGATAG GGACCGAACT	3480
GTCTCACGAC GTTCTGAACC CAGCTCGCGT	3510

(2) INFORMATION FOR SEQ ID NO: 54:

- (i) SEQUENCE CHARACTERISTICS:
- (A) LENGTH: 20986 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 54:

CGGAGAAAAA CATGGCTAAG TCAAACTTG AAAAGTAGA ATCAGTTGTT GGCTGGTTC	60
GTGATAAGAA AATCACAGGC TACCGTATCT CTAAGAAC GAATGCGCGT GAAATGTCTA	120
TCATTGCTCT GGCGCAGGGT CGTGCAAAAG TAAAAATAT TTCATTGAA ACAGCCCTAG	180
GCCTAATTGA TTTCTATGAA AAAATTATG AAAATTGAA AGATTAATCT TTGGATAACG	240
GCGGATTCTT GACCTCAAG TAGTAGAGAT AGAGAACCTG CCTTTTCATT TTGAGGACAG	300
CAAAAAGACT GCACGGTTGA TGCAGCCTT TCTTTTATT TGAGATAGCG TTGAAGGAAC	360
TCTTTGTTTC GGTCTCTTT AGGATTGGTG AAGAGGTCTT CTGGTTTACCT TTCTTCAGCG	420
ATCACCCCT TATCCATAAA GATAACACGG TGAGAGACAT CACGGCAAA TTCCATTTC	480
TGGGTTACGA CAATCATGGT CAAGCCTTCC TGAGCCAGGT CCTGCATGAT TTTGAGGACT	540
TCTCCAACCA TTTCTGGATC GAGAGCTGAT CTTGGTTCAT CAAAGAGAAT AGCGTCCGGA	600
TTCATGGAGA GGGCAGGAGC GATGGCCACA CGTTGTTTT GACCACCTGA GAGTTGTTT	660
GGTTGGCTT GCCAGTAGCG TTCTCCCATG CCGACCTTT CCAGGTTTTC TTTGGCAATC	720
TTTTCAGCTT CTGTGCGTTC CGGTTTGTAG ACAGTTGTCT GAGCGACGAT TGTGTTTCA	780
AGAACATTGA GATTTCAAA GAGGTTAAAG GATTGGAAA CCATCCCCAA CTTTCACGG	840
TATTGCGTGA GGTCAAGGCC TTTTCGAGG ACGTTTGTC CATGATAAAG GATTGTC	900
TCAGTTGGTG TTTCAAGTAG GTTAATGGAG CGTAGGAAAG TCGATTTCC GCTTCCAGAG	960
CTTCCGATGA TAGAGATGAC CTCTCCCTG TGGACAGTGA GTGAAATGTC TTTTAGCACT	1020
TCGTTTGTC CATAGGATT TTTGAGGTGT TTAATTCAA GGATTGCTTG TGTCAATTATT	1080
TCAAATCCTC CGTTGCATT TGGTTAGCAC CTGTAGTGTAA GGTATCCATG TCCATTCTGC	1140

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GCTCGATAAA GCGTAGGATA CGTGTACGG TGAAGGTGAG GACAAAGTAA ATCACGGCGA	1200
TGATTGATAA TGTCTGGAAG TATTGATAGG TTTGTGTTGC CACGGTATTT CCTGAGAAAT	1260
AAAGTTGAC AACAGAGATA ACGTTCAATA CAGATGTATC TTTGATATTG ATGACAAATT	1320
CATTACCAGT TGCAGGTAGG ATGTTACGGA CTACCTGAGG TAGGACAATC TTACGCATGG	1380
TCTGGTTATG GGTCATACCA AGAGCAGTCG CAGCTTCAA TTGTCCTTG TCAACTGCTA	1440
GGATACCAAC ACGGACGATT TCAGTCATGT AGGCACCGGT ATTGATTGAA ACGATGAAGA	1500
TAGCAGCCAG TGTACGGTCA AGGTTGATCC CGAAAGCTTG GGCAGTCCA TAGTAGATAA	1560
CCATCGAATTG AACAAATCATT GGCCTTACAC GGAAAATTTC AATGTAGACA TTGAGAACCC	1620
AGCCGACTAG TTTMTGTTAGG CCGTAAATGA CTTTGTTCAG AGAGAGAGGA GCAGTACCGA	1680
AGACACCAAT GGCAAGTCCA ATAATGAGAC CTATGATGGT TCCGACGATA GAGATTAAGA	1740
GAGTGATACC AGCACCCACGC AAGAGTTGTT GCCAGTTTC AGAAAAGAATT TTAGCAACTT	1800
GGCTAAAGAA ACTACTGCTA GTCTCTTCAG TTGTTGAGC TTCGGCAGGT TGTTCTTGA	1860
TCATACGATC CATCAAGGCA ACTTGGTCAT CTTTGAAAT GGTTCAATG CTGGCATTGA	1920
TTTGGCTAAT ACGATTGTCA TTTTACGAA GCCCGATAGC GATAGCTGTA TCTTCTTCCC	1980
CAGTTTGAA ACCAGGTTCT ACTTGAATCA TCTTGAACTT AGAGTTGCA GCTTCAGCAG	2040
TCAGTGCTTC TGGACGTTCA GAAACATAAG CATCAATGAC ACCAGCCTCA AGAGCTTGTC	2100
CCATTTGAGC GAAGCTCTCC ATGGCTGTTT CTTTTTGTAGC ACCTGGGATT TGTGCAATCA	2160
AGTTATAAAAG GTAGACCCCT TGTTGAGAAG TGATTTTGC ACCGTTAAAG TCATCCAAAG	2220
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GTCTTACGG TAGGTATTGG TATTAACGGA GATCCTGGTA CGGATGAAGC AACGGCTACT	16320
AGTTTATGC AAAGTATTTC TAGTAAACCT GAAAACATAA CCAATGTTAC TGACACGACA	16380
AAAATATTGG AACAGTTGAA TCGTTATTTC CACACCATCG TAACTGAAAA GAAATCAATT	16440
GAGAATGGTA CGATTACAGA TCCGATGGGT GAGTTAATTG ATTTGCAATT GGGCACAGAT	16500
GGAAGATTTG ATCCACGAGA TTACACTTTA ACTGCAAACG ATGGTAGTCG CTTGGAGAAT	16560
GGACAAGCTG TAGGTGGTCC ACAAAATGAT GGTGGTTGT TAAAAAATGC AAAAGTGCTC	16620
TATGATACGA CTGAGAAAAG GATTCGTGTA ACAGGTCTGT ACCTTGGAAC CGATGAAAAA	16680
GTTACGTTGA CCTACAATGT TCGTTTGAT GATGAGTTG TAAGCAATAA ATTTTATGAT	16740
ACCAATGGTC GAACAACCTT ACATCCTAAG GAAGTAGAAC AGAACACAGT CGCGCAGTTC	16800
CCGATTCTTA AGATCGTGA TGTGCGGAAG TATCCAGAAA TCACAATTTC AAAAGAGAAA	16860
AAACTGGTG ACATGGAGTT TATTAAGGTC AAAAAAAATG ATAAAAAAACC ACTGAGAGGT	16920
GCGGTCTTTA GTCTTCAAAA ACAACATCCG GATTATCCAG ATATTTATGG AGCTATTGAT	16980
CAAAATGGCA CTTATCAAAA TGTGAGAACCA GGTGAAGATG GTAAGTTGAC CTTTAAAAAT	17040

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CTGTCAGATG GGAAATATCG ATTATTTGAA AATTCTGAAC CAGCTGGTTA TAAACCCGT	17100
CAAAATAAGC CTATCGTGC CTTCCAAATA GTAAATGGAG AAGTCAGAGA TGTGACTTCA	17160
ATCGTCCAC AAGATATACC AGCGGGTTAC GAGTTTACGA ATGATAAGCA CTATATTACC	17220
AATGAACCTA TTCCCTCCAAA GAGAGAAATAT CCTCGAACTG GTGGTATCGG AATGTTGCCA	17280
TTCTATCTGA TAGGTTGCAT GATGATGGGA GGAGTTCTAT TATACACACG GAAACATCCG	17340
TAAAGTGTAG AAATGATAAT ATCTATGTT TGAAACGATAC TTTTAAGAAG TAGCACTCAA	17400
GAAGAGATTT AAGTTTACTT GGTGAAACCT GTTTTATTG TAAGTAAACT ATCATTGAAA	17460
GGGGAGATGT TTTCGAAAC TTGCACAGAA AAAGGATTAT TATTGTCATG TGTAATTCAT	17520
TACATTGCTC ACAGTTGATT TTAAGAGATA TGAATAAGGA GAAATCATGA AATCAATCAA	17580
CAAATTTTA ACAATGCTTG CTGCCTTATT ACTGACAGCG AGTAGCCTGT TTTCAGCTGC	17640
AACAGTTTT GC GGCTGGGA CGACAACAAAC ATCTGTTACC GTTCATAAAC TATTGGCAC	17700
AGATGGGGAT ATGGATAAAA TTGCAAATGA GTTAGAAACA GGTAACATATG CTGGTAATAA	17760
AGTGGGTGTT CTACCTGCAA ATGCAAAAGA AATTGCCGGT GTTATGTTCG TTTGGACAAA	17820
TACTAATAAT GAAATTATTG ATGAAAATGG CCAAACCTA GGAGTGAATA TTGATCCACA	17880
AAACATTAAA CTCTCAGGGG CAATGCCGGC AACTGCAATG AAAAAATTAA CAGAAGCTGA	17940
AGGAGCTAAA TTAAACACGG CAAATTACCG AGCTGCTAAC TATAAAATTG ATGAAATTCA	18000
CAGTTTATCA ACTTATGTCG GTGAAGATGG AGCAACCTA ACAGGTTCTA AAGCAGTTCC	18060
AATTGAAATT GAATTACCAT TGAAACGATGT TGTGGATGCG CATGTGTATC CAAAAAATAC	18120
AGAAGCAAAG CCAAAATTG ATAAAGATT CAAAGGTAAA CCAAATCCAG ATACACCACG	18180
TGTAGATAAA GATACACCTG TGAAACCACCA AGTTGGAGAT GTTGTAGAGT ACGAAATTGT	18240
TACAAAAATT CCAGCACTTG CTAATTATGC AACAGCAAAC TGGAGCGATA GAATGACTGA	18300
AGGTTGGCA TTCAACAAAG GTACAGTGAA AGTAACGTT GATGATGTTG CACTTGAAGC	18360
AGGTGATTAT GCTCTAACAG AAGTAGCAAC TGGTTTGAT TTGAAATTAA CAGATGCTGG	18420
TTTAGCTAAA GTGAATGACC AAAACGCTGA AAAACTGTG AAAACTACTT ATTGGCAAC	18480
ATTGAATGAC AAAGCAATTG TAGAAGTACC AGAATCTAAT GATGTAACAT TTAACATATGG	18540
TAATAATCCA GATCACGGGA ATACTCCAAA GCCGAATAAG CCAAATGAAA ACGGCGATT	18600
GACATTGACC AAGACATGGG TTGATGCTAC AGTGCACCA ATTCCGGCTG GAGCTGAAGC	18660
AACGTTCGAT TTGGTTAATG CTCAGACTGG TAAAGTTGTA CAAACTGTAA CTTTGACAAC	18720
AGACAAAAAT ACAGTTACTG TTAACGGATT GGATAAAAAT ACAGAAATATA AATTGCTGA	18780
ACGTAGTATA AAAGGGTATT CAGCAGATTA TCAAGAAATC ACTACAGCTG GAGAAATTGC	18840

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TGTCAAGAAC	TGGAAAGACG	AAAATCCAA	ACCACATTGAT	CCAACAGAGC	CAAAAGTTGT	18900
TACATATGGT	AAAAAGTTG	TCAAAGTTA	TGATAAAGAT	AATCGTTAG	CTGGGGCACA	18960
ATTTGTAATT	GCAAATGCTG	ATAATGCTGG	TCAATATTAA	GCACGTAAG	CAGATAAAAGT	19020
GAGTCAGAA	GAGAAGCAGT	TGGTTGTTAC	ACAAAGGAT	GCTTTAGATA	GAGCAGTTGC	19080
TGCTTATAAC	GCTCTTACTG	CACAACAACA	AACTCAGCAA	GAAAAAGAGA	AAGTTGACAA	19140
AGCTCAAGCT	GCTTATAATG	CTGCTGTGAT	TGCTGCCAAC	AATGCATTG	AATGGGTGCC	19200
AGATAAGGAC	AATGAAAATG	TTGTGAAATT	AGTTTCTGAT	GCACAAGTC	GCTTTGAAAT	19260
TACAGGCCIT	CTTGAGGTA	CATATTACTT	AGAAGAAACA	AAACAGCCTG	CTCGTTATGC	19320
ATTACTAACT	AGCCGTCAGA	AATTGAAAGT	CACTGCAACT	TCTTATTTCAG	CGACTGGACAA	19380
AGGCATTGAG	TATACTGCTG	GTTCAAGGTA	AGATGACGCT	ACAAAAGTAG	TCAACAAAAAA	19440
AATCACTATC	CCACAAACGG	GTGGTATTGG	TACAATTATC	TTTGCCTGAG	CGGGGGCTGC	19500
GATTATGGGT	ATTGCAGTGT	ACGCATATGT	AAAAAACAAAC	AAAGATGAGG	ATCAACTTGC	19560
TTAAGTAAGA	GAGAAAGGAG	CCATTGATGA	CAATGCAGAA	AATGCAGAAA	ATGATTAGTC	19620
GTATCTCTT	TGTTATGGCT	CTGTTGTTTT	CTCTTGTATG	GGGTGCACAT	GCAGTCCAAG	19680
CGCAAGAAGA	TCACACGTTG	GTCTTGCAT	TGGAGAACTA	TCAGGAGGTG	GTAGTCAAT	19740
TGCCATCTCG	TGATGGTCAT	CGGTTGCAAG	TATGGAAGTT	GGATGATTG	TATTCCTATG	19800
ATGATEGGGT	GCAAAATGTA	AGAGACTTGC	ATTCGTGGGA	TGAGAATAAA	CTTTCTTCTT	19860
TCAAAAGAC	TTCGTTGAG	ATGACCTTCC	TTGAGAATCA	GATTGAAGTA	TCTCATATT	19920
CAAATGGTCT	TTACTATGTT	CGCTCTATTA	TCCAGACGGA	TGCGGTTCT	TATCCAGCTG	19980
AAATTCTTTT	TGAAATGACA	GATCAAACGG	TAGAGCCTTT	GGTCATTGTA	GCGAAAAAAA	20040
CAGATACAAT	GACACAAAG	GTGAAGCTGA	AAAGGTGGA	TCAAGACCAC	AATCGCTTGG	20100
AGGGTGTGCG	CTTTAAATTG	GTATCAGTAG	CAAGAGATGT	TTCTGAAAAA	GAGGTTCCCT	20160
TGATTGGAGA	ATACCGTTAC	AGTTCTCTG	GTCAAGTAGG	GAGAACTCTC	TATACTGATA	20220
AAAATGGAGA	GATTGGTG	ACAAATCTTC	CTCTTGGAA	CTATCGTTTC	AAGGAGGTGG	20280
AGCCACTGGC	AGGCTATGCT	GTTACGACGC	TGGATACGGA	TGTCCAGCTG	GTAGATCATC	20340
AGCTGGTGAC	GATTACGGTT	GTCAATCAGA	AATTACCACG	TGGCAATGTT	GAATTATGAA	20400
AGGTGGATGG	TCGGACCAAT	ACCTCTCTTC	AAGGGCAAT	GTTCAAAGTC	ATGAAAGAAG	20460
AAAGCGGACA	CTATACTCCT	GTTCTTCAAA	ATGGTAAGGA	AGTAGTTGTA	ACATCAGGGA	20520
AAGATGGTCG	TTTCCGAGTG	GAAGGTCTAG	AGTATGGAC	ATACTATTTA	TGGGAGCTCC	20580

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AAGCTCCAAC TGGTTATGTT CAATTAAACAT CGCCTGTTTC CTTTACAATC GGGAAAGATA	20640
CTCGTAAGGA ACTGGTAACA GTGGTTAAAA ATAACAAGCG ACCACGGATT GATGTGCCAG	20700
ATACAGGGGA AGAAACCCCTT GTATATCTTG ATGCTTGTG CCATTTGTT GTTTGGTAGT	20760
GGTTATTGTC TTACGAAAAA ACCAATAAAC TGATATTCAA TGTACATCAT TATGAATAGG	20820
ATAGCAGGCT GAAGGGAAGA CCAGAGTACT CTGAGGTGAT GTTAATCAGG AATCATGGTG	20880
ATGTGGCATG AATCATCAAT AACCGATATG AGGCTGGCA GATTGTGCCA GCCTCATTGT	20940
GGGTTATTGTT TTGAAAACG ATAGGACTGG TCTGGTAATC ATTTTA	20986

(2) INFORMATION FOR SEQ ID NO: 55:

- (i) SEQUENCE CHARACTERISTICS:
- (A) LENGTH: 21040 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 55:

CCAGCAAAA AGCCATCCGA AGATGACTTT TTTGCTATTT AATTCTGTA TAAGTTACTT	60
CCAAGCCACG CTTAACAGCT GGACGATTGG CAATTTTTC TGCCCATTTC ACTAGATTTT	120
GATAACTTGA GGCATCCAAG AATTTCAGAC AACCTTGGTA AAGATTCCCT TGAACACTAATC	180
GTCCATACCA AGACCAGATA GCAATATCTG CAATCGTATA GTCATTGCCT GCAATATAAG	240
GTTPCTGAGC CAATTCCTTA TCCAATAAAAT CCAACTGGCG TTTCACTTCC ATCGTAAAAC	300
GGTTAAATAGG ATATTCCAAT TTTTCAGGAG CATAATTGAA GAAATGTCCA AATCCCCCAC	360
CTAGAAAAGG TGCTGCACCT GCTTGCAGA ATAGCCAATT CAAAACCTCT ACCTTTTCCA	420
CAGGATTACT TGGTAAAAG GCTCCAAATT TCTCAGCAAG GTAAAAGAAGA ATATGAGCAG	480
ACTCAAAGAC TCTTACGTTT TCAGTACCTG ACTGGTCCAA TAAGGCTGGA ATCTTGGAAAT	540
TTGGATTGAG CTTCACAAAG TCTGATCCGA ATTGATCCCC ATCCATGATA GCAATCTTAT	600
ACAAGTCGTA AGCCGCTTCC TTAAAACCAG CTTCTAGTAA TTCTTCCAAT AAGATAGTAA	660
CCTTCACACC ATTTGGTGTG CCCAGTGAAT AAAGCTGAAA AGCTTGTCT CCTTTGGCA	720
AGTTTTGTTC GAAACGGGCA CCTGCTGTTG GTCTGTTAG CCCCCGAAAA GCTCCTTGAT	780
TACTAGCTTC ATCCGCCAT ACGGTGGTA ATTGATATGC TGACATCCGA AACCTCCCTT	840
AAATCCGATT CTTGCAAAA CCGAGTTGC GTTGAATAAA CTTAACGATT TCGACGATGAA	900
TAATCATTGA GAAGCTTCCA GCCATAACAA TTCCCCATTG TGACAAAGTCT AGTTGGTTA	960
CGTGGAAAGAT TCCTCAAGC GGTTCTACAA CGATTGTTGC CATGAGAAGG ATAAAGGATA	1020

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CCAAGATGGA CCAGTTAAAG GTCTTAGACT TGAATGGCC AACTGTCAAG ATGGATTGGT	1080
AGACACACTT GACATTGTAG GCATGGAAGA GCTGAATCAA ACCAAGGGTT GCAAAGGCCA	1140
TCGTTAGGGC ATCTGCATGA ATAGCATGAT TGTCACCCAC ATGAACTGGG TAAGCAATCG	1200
CAAGGCCATA AACACTCATA ACAAGAGCTG CTTGGAGTAC ACCTTGATAA ATGATAGAAC	1260
TCAAAACACC ACCTGAGAAG AAGCTTGCT TCCGTCCACG TGTTTATGA TTCATGACAC	1320
CAGGTTCCGC AGGTTCAACA CCAAGAGCGA TAGCTGGAA GGTATCCGTT ACCAAGTTGA	1380
TCCACAAAAG ATGAACCGGC TGTAAGACAT CCCAACAAA CAAGGTTGAT AGGAAGATGG	1440
TTAATCTTC AGCAGTATTA GCAGAAAGTA GGTAAGTAAAGTCTTTGA ATGTTTGAGA	1500
AGACCTTACG TCCTTCTTCC ACTGCGACGA TAATAGTCGC AAAGTTATCA TCTGCAAGAA	1560
TCATATCAGA AGCCCCCTTA GAAACCTCTG TACCAAGTATGAT TCCCACATCCG ATACCGATAT	1620
CGGCTGTTT CAGAGCTGGC GCGTCATTGA CACCGTCACC TGTCATGGCA ACGACTTTAC	1680
CTTGTTTTG CCAAGCCTTG ACGATACGAA CCTTGTGTTG TGAGACACA CGGGCATAAA	1740
CAGAGTATTG ACCAACGACT TTTCAAAATT CTTCATCTGA CAGTTCAATTG AGTTCAAGCAC	1800
CAGTTAAAAC GTGACCTTCT GTATCGTTG CGTCAATGAT TCCCAAAACGT TTGGCAATGG	1860
CTTCCGCTGT GTCTGGTGG TCACCTGTAA TCATAATTGG ACGGATTCCC GCTTCCTTAG	1920
CCACACGAAC AGCCTCAGCG GCTTCAGGAC GTTCAGGTC AATCATCCCA ATCAAACAG	1980
TAAAAATTAA ATCATTTCA AGCTCTTCAG AAGTGAGATT TTCTGGAATA CTATCGATAA	2040
TCTTATAAGC ACCTGCAAGG ACACGCAAGG CTTGATGAGC CATTTCAGAA TTGTTTGTAC	2100
GAATGAGATT TGTAACCTTC TCATCAATCG GAGCAATATC CCCAGCCTTA TCACGAAGAA	2160
GACAACGTTT TAAGAGTTGG TCTGGCGCAC CCTTGACTGC TACAAGGAAA CGACCACATCTG	2220
GCAATGGGTG AACTGTTGAC ATGAGCTTAC GGTCAAGAGTC AAATGGCAAT TCAGCTACAC	2280
GAGGATATTCTCTAAGAAA CCTTGACAT CATAGCCCTT GTCCAAGGCA TATTGGATAA	2340
AGGCTGTTTC GGTTGGGTCA CCAATCAAGT TACCTTCCAC ATCGATTTTC GTATCATTGG	2400
CCAAGACAAC TGAACGAAGT AGTGGCATTT CAAGACCTAG TTCAATATCA TCAGCTGAGT	2460
CATGTAGAAC CGCATCGTAG AAGACTTTTT CGACTGTCAT CTTGTTCATCA GTCAGCGTAC	2520
CAGTCTTATC AGAAGCGATG ATTCAGTTG AACCAAGTGT TTCAACTGCT GGCAACTTAC	2580
GAACGATGGA ATGTCGTTG GCCAAAACCTT GAGTACCAAG AGAAAGAACG ATGGTAACGA	2640
TAGCAGGAAG TCCTTCTGGA ATGGCTGCAA CGGCAAGGGC AACAGAAGTC AACAAACTCAC	2700
CAAGTGGATT TTTCCTTGA ATGAAGACAC CCACTACAAA AGTAACAAGG GCAATGACCA	2760

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AGATAGCATA GGTCAAGACC TTAGAAAGGT TGTTCAAATT TTGTTTGACT GGTGTATCAG	2820
TCTCATCCGC ATCTTGAAGC ATACCAAGCAA TATGCCAAC TTCAGTGTAC ATACCTGTAT	2880
TGACAACAAAC ACCCATCCCA CGACCATAGG TTACGTTGA GTTTTGGAAAG GCCATGTTGA	2940
CACGGTCACC AATACCAGCA TCTGTCGAA GCTCGACTGA CAAGTCTTT TCGACTGGTA	3000
CAGATTCAACC TGTCAAGGCT GCTTCTCAA TTTTAAGAGA GTTGGCTTCT ATCAAACGTA	3060
GGTCCGCTGG TACCAAGCTCA CCTGCTTCAA GGGCAACGAT ATCGCCTGGT ACCAATTCTT	3120
TAGAGTCAAT CTCTGCCATG TGTCATCAC GAAGAACGCG GGCAACTGGA CTAGACATGG	3180
ATTTGAGGGC TTCAATAGCT TCTTCAGCTT TTCCCTCTTG GTAAACACCA AAGGCAGCGT	3240
TGATGATAAC CACAGCTAGG ATGATAATGG CATCTGCGAT ATCTTCCCCA CCAGAACGTA	3300
CGACTGACAA GATTGctGCC GCAACTAGGA TGATAATCAT CAAATCCTTA AATTGCTCGA	3360
TGAATTGAC CAAGATTGAT CGTTTCTCGC CTTCCTCGAG TTCATTGTGC CCAAATTCCG	3420
CAAGGCCTT TTCCGCCTCA CTTGATGACA AACCTTGCTC GGTGCGCATCC ACAGCCTGCA	3480
AGACCTCTTC AGGGCTCTGA GTATAAAACG CTTGGCGTTT TTGTTCTTT GACATGTGTC	3540
TCTCCCTGA CATTGTCGCA AAAACAGACT CTCTTCTGT CATAGCTTT CACGACAAAC	3600
AAAAAGAAC CTGTTAATCA TAACAAGTCT CGCTGTTAA GATAGGGCCG GAAAGCATAAC	3660
TTTTCAGCAT AAAATTCCGA ATGACGACAC TATCACAGGT TTCTGCCAGC TACTCCCTTG	3720
AGTAGTACCA TTATACAAA TTTTGGGAG TTTTCAAAGA GTAAAAACTG CCTTATTGAA	3780
ATTTTCCTT GAAAACAGT ATAATGGTAG AATGCTATGT GACTAGAAAG GAAGTTGAAT	3840
GAAGCAATCT ATCTCAAATC TCAAGTTAGC TGAGCGTGGA GCCATTATCA GTATTTCGAC	3900
CTATTTGATC TTGTCGTCAG CCAAATTAGC AGCTGGTCAT CTCCtTCATT CATCCAGTT	3960
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GATTGGATG GCGGCCACC TGCAGACCGT GACCACCGTT TTGGTCATTG GAAGATTGAA	4080
GATTTGGCAA GCTTGATCAC TTCTATCATC ATGTTCTATG TCGGTTTCGA TGTTCTAAGA	4140
GATACCATTG AAAAGATTCT CAGTCGGAA GAAACGGTC A TTGATCCTCT TGGTGCAACT	4200
CTAGGAATCA TTTCTGCAGC GATTATGTTT GTGGTCTATC TCTACAATAC TCGCCTCAGT	4260
AAGAAATCCA ACTCCAATGC GCTGAAGGCA GCTGCTAAGG ACAATCTTC TGACGCTGTT	4320
ACCTCACTTG GAACCGCCAT TGCCATCCTA GCTAGTAGTT TCAATTATCC GATTGTGGAT	4380
AAACTGGTTG CTATCATCAT CACTTTCTTT ATCTTGAAGA CTGCTATGA TATCTTCATC	4440
GAGTCCTCCT TTAGTCTTTC AGATGGCTTT GACGACCGCC TGCTCGAGGA CTACCAAAAG	4500
GCTATCATGG AAATTCCCAA AATCAGCAAG GTCAAATCGC AAAGAGGTCG CACCTACGGT	4560

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AGCAACATCT ACCTGGATAT TACACTAGAG ATGAATCCTG ACTTGTCTGT	TTTGAAAGC 4620
CATGAAATCG CGGATCAGGT CGAGTCTATG CTGGAGGAGC GTTTGGCGT	CTTTGATACC 4680
GATGTCATA TCGAACCCAGC ACCTATCCCT GAGGATGAAA TTTAGACAA	TGTCTATAAA 4740
AAATTGCTTA TGCCTGAACA ATTGATTGAC CAAGGAAACC AACTAGAAGA	ACTCTTGACT 4800
GATGATTTG TCTATATTG CCAAGATGGA GAGCAGATGG ATAAAGAGGC	TTATAAGACC 4860
AAAAAAAGAGT TAAATTCTGC TATCAAGGAC ATTCAAATTA CTTCCATCAG	TCAAAAAACC 4920
AAACTCATCT GCTATGAGTT AGATGGTATC ATCCATACCA GTATCTGGCG	TCGCCACGAA 4980
ACCTGGAAA ATATCTTCA TCAAGAAACC AAAAAGAAT AGAGAAATCC	TTTCATGAGA 5040
CGGGATTTT CTATCTTTT ATACTCAATA AAAATCAAAG TGCAAATTAG	GAAGCCGGTC 5100
ACAGGCTGTA CTTGAGTCGG CAATGTGAAG CCGACATAGT TTGCACTTTG	ATTTTCGAAT 5160
AGTCTTAACt ATCAAATTCA CTGAGATACT CATAGCGTTC GTATTTTCA	AGGAGTGCTT 5220
CATTTTCTC ATCCAATTCT TTTTGGAGAG TAGCCAGCTT ACCAAAGTCA	GAGCCGTTAG 5280
CCTGCATTTG CTCTTCATA GCAGCGATAC GTTTTCCAA GGTTTCAATA	TCACCTTCAA 5340
TACTTGCCC CTCCGCTTT TCTTGGTAGG TCATGCGTTT CTTGTCTTCT	CGAACCTTGA 5400
CCACTTTTC CTTTTCGGCC TTTTGCACTT GATTGGCCAT ATCTGTTCA	AAAGCTTTT 5460
CATCAAAGATA GTCGGTGTA TGACCAAAGA AAGGACGAAT CTTGCCATCC	TCAAAAGCGA 5520
GAACTTGGT CGCTACCTTA TCCAAGAAAT AGCGGTCGTG ACTGACTGTT	AAAACGGGAC 5580
CTGCAAAACC TTGCAAGAAA TTCTCTAAGA CTGTCAAAGT TGCAATATCT	AGGTCAATTGG 5640
TTGGCTCGTC TAAAAGAAGA ACATTTGGTT TTTCCAAAG CAGTTGAGG	AGATAAAAGAC 5700
GTTTTTCTC ACCCCCTGAC AATTCTCAA TCAAAGTCCC ATGCGTCGAA	CGTGGGAAGA 5760
GGAAATTGCTC CAGCAACTCA GCGATGGAAG TCGTAGAACC ACCACTGGTC	TTGACCTCCT 5820
CTGCCACTTC CTGCAGGTA TTGATCACAC GCTTGCTTTG ATCCAAACCC	TCAATTGTT 5880
GAGAGAAATA GGCGATCGCA ACAGTTTCCC CAATCACAAC TTGTCCTGCT	GTCGGCTCAA 5940
GACTTCCCTGC AATCAGGTTA AGTAGGGTTG ATTTTCCAAC ACCATTGTCC	CCAACAAATTG 6000
CAATACGGTC TTTAGCCTGA ACTAAGAGAT TAAAATTG CAAAATGGGC	TTATTTTCAT 6060
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GATTGATACG AGCTTGTGTC TTGGTCGCAC GCGCCTGCGG TTGTCTGCGC	ATCCAGGCCA 6240
ATTCTMTGTTT GTAGAGTTGT TCTTTTTGT GAAGAAGAGC CGCGTCGCGC	TCATCCTGTT 6300

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CCGCCTTAG GCGAACATAG TCCTGTAAT TTCCCTGGTA CTCGGTCAAG CCTGCACGAT	6360
CCAACTCGAA AATCCGTGTT GACAAGCGT CTAAGAAATA ACGATCGTGA GTGATAAAAAA	6420
GGACGGCTT CTTAGAATT TTCAAAAAGA GGTCAGCCA CTCATAATC GCAATATCCA	6480
GATGGTTGGT CGGCTCATCC AAAAGCAAGA GGTCGTGGTT GCCAAGTAAG ACTTGTGCCA	6540
ACTGTACCCG TCTTCTCAGA CCACCTGACA ATTCCCCAAC AGGAGTAGAT AAGTCTTGAA	6600
TGCCAATTT GCTAAGAACG GTCTGACCT GACTTTCGAT TTCCCAAGCT TGGAGAGAGT	6660
CCATCTCTGC CATGACACGT TCCAAACGCG CCTGCTTGTC CTCACTATAG TCGAGCATAA	6720
TCAATTCTATA CTCACGAATG AGCTGGATTT CCTTGAGTTC ACTAGATAGA ACCGTATCCA	6780
AAACTGTCTT TCTATCATCA AAATCAGGAT CCTGAGTCAA GTAACCAATC TGGTAATCAT	6840
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AAAGGGTGGT CTTGCCAGTC CCATTGACAC CGATTAAACC AATTCTGTCT AAGTCATGGA	6960
TAATAAAGGA AATATCCCTA AAAACGGTCT TGTCACCAAC GGATTTACTT AGTTTTCAA	7020
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CAATTCTCCA TCGACAATGG CAAACTCAAT CTCTGTTAAA ATCTCTCCCAGTCTGGGCC	7140
TGGCTGATAG CCATATTCTTGATG TGATCAAAAT ACCGCCATTA ATCTGAATCT CTTTCTTGTC	7200
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AGCTTGACGA AGATTTCAAG CCTGTAAAAG CAAATCTATG TCAAAGCGAT AACAAATCTCG	7320
CTTGCTCAAT TCTCCATTTCAG CACCGAGC CAAATAATC AGCAAACTCTT GAACCTTGCTT	7380
GGCAAACCTGG CGTGAGGTCT TCCAAGATTT CAAAATGAC TGCGCATTTT CAATCTCCAA	7440
AGCCCCATAGT AAAGCCGCC AGGCTTGTTC AGAGGATTCA AAAGTAAAT CAGTCTCCAA	7500
ATCAAACAGT CTGTTGAGCT TGTCCTGGCT AGATGCCATA TCAGGGAGAT AGTCATAAGC	7560
TTGACTCTCA ATCATGGAAG CCAAGCCCCCT TCTCCAAAAT GGAGCCAGCA AGAGTTTATC	7620
AAACTCGACG AAGGTACGCT CTACAGAAAT TTTCTCCAAA AGCGGGGTCA AGGTCTTCAT	7680
AGCTTTAAAT GTTCTGGCT CAAGTGCAGAA ACCAAGACTA GCCTGAAAC GGAAACCACG	7740
CATAATCCGT AAAGCATCTT CGTGAAACG CTCACTAGCC ACTCCAACG CTCGCAAGAC	7800
TTGCTTTCC AAATCTCTA AACCAGGAA CAAGTCAACG ATTTCTCTG TCTCATCCAA	7860
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GGAAACCGCA CTGGGCTGCA GATAAGTCCAC ATAGACATCC TCTGTCCGAA AGGTTGTTAC	7980
CTCATACTCC TCATCCCCAT CTAAGACCAA GACGGTTCCA TGCTCGATTC CGATATCGGC	8040
TGTTCCGGAA AAAATCTGCT TGGTCTCTTC TGGATAAGAA GACGTCGCAA TATCCACATC	8100

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GTGGATAGGG CTATGGAGAA GGGCATCTCG AACAGAGCCC CCAACAAAAT AAGCCTAAA	8160
GCCTGCTTCT TTAATTTTTT CTAATACTGG TAAAGCCTTC TGAAATTCAAG AAGGCATTTG	8220
CGTTAATCTC ATAATAAGTG TTCTAATCCA TAGACAAGCT CATGACGCTT GACAACCTCT	8280
TTAATTCCCA AATTGACTCC TGTCACTGAAG GAGATGCGAT CATAGGAGTC ATGACGGAGG	8340
GTCAACCCCTT CTCCCTGATT GCCAAAGATG ACTTCCTGAT GAGCTACCAA GCCTGGAAA	8400
CGAACTGAGT GGATGCGCAT ACCATCAAAG TCAGCACCAAC GAGCACCAGC AATCAGCTCT	8460
TCCTCATCTG CTGCACCTTG CTGAATTGAC TCTCGAACCT CTGCCATCAA CTCAGCTGTT	8520
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GCAAAGTTAG GGGCAATCAG GCCACCCAAG TCTTGGGCAC GAGAAAATTC TTTTAGCTCT	8700
GCAATTCTT CACTCGTGA ACCAGTCGTT CCAACTACTG GAGCAAAGCC ATTTTCAAGA	8760
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GAAGTGTCTA CTTCTTGGAA GAACTATCTT TTTCACACAG GGTTCCAGGC GTGTTCAATT	9120
ATCAAGATAAC AAAGGACCTT AGCTGCCTCT GAAAAATAGG GAATGGCACT GACTTTCCAC	9180
GAAAGGCAAG ACAGGCATCT TTTTCAAGA GGCAGGTAGT CGGTGTTCAA TTTCTAAGAT	9240
ACAAGGCATC TTAACTAGCC TAGAAGCGCC AACTAAATCA CTGGAATATA ACCCAGAGCA	9300
ATACTCCTG CTCCCTAGGTG CGTTCCAATG ACACTACCAA ATGTAGCAAG TGAAACATCC	9360
GAACCCAAGC CAAAATCAAG CAAGTGCTGA CGCAATTCTT CAGCCTTTTC AGGAGCATT	9420
CCATGAATGA CAATGACCCG GTATTGACCT GAAGCCGTTG TTCCCTGAT AATTTCAATT	9480
AAGCGCTTGG TGGCCTTCTT TTCAGTACGA ACTTTTTCTG AAACATTCAAT CACACCTG	9540
TCGTTAAAAT AAAGGATTGG CTTAATGCTA AGCAAATTGC CCAAATGGC AGCCCCATT	9600
GAAAGGCGTC CACCTTTAC CAAATGATCC AAGTCATCTA CCATGATAAA GGCTGACGTA	9660
CGGCTGATTT GAATGGCTAG CTTATCCTGA ATGCTGGCAA AATCATCGCC CTGATCACGC	9720
CAATTTAAAGA CGCTTCAAC CATGATGCCT AGGGGAGCAC TTGTAATCAA AGTGTCTGGG	9780
AAAGCAATGG TTAAGCCCTC ATAGTCATCG ACCATATACT GGATATTTG GTAAAAACCT	9840

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GAAATTCCAG	AAGATAGGAA	AAGCCCCAAG
GCATGTGTAT	AGCCTTGTTC	TTTGAGCGAA
9900		
GTAAAGATCT	CATCTAACCT	GGCAATACTT
GGTTGACTGG	TCTTAGGCAA	TTCAGAAGCC
9960		
TGAGCCATT	TTTGGTAAA	TTCCCTCAGCA
GACAGATTGA	TGCCCCTCGAC	ATATTCCCTCA
10020		
CCATCAATAT	TGACAGGAAT	ATCCAAGACA
AACAAGTCTT	CTCTTGC	GA
10080		
CTGAGATAAG	CAGAGGAATC	TGTGAAACA
GCTAATTCA	TATTAGAACT	CCAAATTAAT
10140		
TCCGGTAAG	TCTAATGCAA	TTTCAGTCAC
TC	TCGTAAGTC	AAACGATTGA
GCATGTCAA	10200	
ACATGGACGA	GCCAAAGGTTT	CCACCTCTTC
TTGGTTCAAT	TCAC	TTGGTT
CATTGACAAT	10260	
ACGGCCATCG	ATATGGTTA	CTTGTGAGAT
TGTTCCACTA	ATGACAAACT	TATCAAATAC
10320		
AATCATAAAG	CTCAAGATGA	CAATCAAGGA
AGTCAC	TTGA	TTGGT
GGT	10380	
CAATTGGAAA	TTCACATCCA	CCTTGGTTTC
AGGAGCTCCA	TTTTCATTTT	CCCATTCAA
10440		
ATTACCGCGCA	TCAAAATGAT	ACTGACTAAC
AAATCTTGT	TCAC	TTAA
GATTCA	10500	
TTTCTCCATC	GGCTACAATA	TTATAAGCTA
TTGTAC	TTTCTTATT	TTTCATCTAGT
10560		
TTTCTAGGAT	TTAGTCATC	CCAATTCAG
CACGAACTAC	ATCTGTGATG	GTATCAACAT
10620		
AGTAGTTTAC	TTCTCTGT	GTAGGCC
CTGCC	10680	
CACTTGGACG	AACAAGGATA	CGGCCGTTCC
CCGCC	10740	
TGGCCTTGAT	AGCTGGCACT	TCCATGGCCT
TTTCC	10800	
TAAC	TTTCTTCAT	TTTCTTCA
AGC	10860	
TTTCTTCA	TTG	TTG
TAAC	10920	
AATCCATCAA	GATAACGTGA	CCAGACTGTT
CAC	10980	
TTTCTTCAC	AACGTAGCGG	TCACCAACTG
CA	11040	
TCAAGGCCTT	GTG	AAAC
AGG	11100	
AT	TTCAAGAAAGG	TATTTCCGA
TG	11160	
TGTCACCATT	CTCATCAACA	GCAATCAAGC
GGT	11220	
TAGCTGACCC	ACTTTCTT	ACCAC
TG	11280	
TAAGGTTGAT	GT	GG
GT	11340	
CTGCAAAGAT	TTGACGGGCA	CTGGT
AG	11400	
TCAT	TTG	GT
AG	11460	
TGGATAATCT	AC	AAATTC
CTA	11520	
AGC	TTG	GT
AG	11580	
GCCAAAGAAC	TTG	GGT
TG	11640	
TATCAAGGGC	TGG	GTG
TG		
CTAGCAGAAA	CTAG	GTG
TCATGACACC	TCAT	GTG

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GGCACTTGCT CCTTCAGTTT CAACCAAGTA AGCTACTGCT GGTGTTGCAA GGACACCAAG	11700
TTTGTATACG TGAATCCCTA CTGAAAGGAG ACCTGCCACC AAGGCCATT CCAACATTTC	11760
CCCTGAAATA CGTGTGTCAC GTCTCACAAA GACTTCGGC GCTTCCGTTT CATGTTGACT	11820
AAGAACATAG CCTCCAAAC GTCTAGTTT AAAGGCTAAT TCTGGTGTAA GTTCTAGGTT	11880
AGCTTCTCCA CGGACTCCAT CAGTCCAAA ATATTTACCC ATTGTTATAA AATCCTTTTC	11940
TATTTTAAAT TCGTTTTGA ACTAGTTGCT TTCGTTGACG AAGATGTCTC CGATGAAC TG	12000
CTTGTACTTG AATTTGATGT GCTTGAACCTT GGTGCTACTG GTTGTGTTAGT CACCTTCATT	12060
ATTGTATCAA ACGGAGTGTAACTGCCGGT AAGACAACAC CATTGCCGTC GATTGCCCTGC	12120
AAAGGTACTG AACCACTGTAA TTACACTGTT ATACGTTCGC TAGTGGCAA AACAGCGATA	12180
ATCTTATCAA TTCTATCCAA TGTCTCTTGG TCACTCGTAA TAGACACTTC TTTATCTGAC	12240
ACCATGACAT TTTCAATTG TACCCGACTA TCAATTGAC TAGGGTCAAT CTCTGGTACA	12300
ATCTTACCT TATCCTCTG AGCCTCTTA CCAATCTTGA CTGTAATT TTGCGGAGTC	12360
GCCACAGCGG TCAGCCCATT GGGTAAATCT TCAATGCTCA AAGGAACTTC AATCGTTCCA	12420
ACACCGGCAT CTGTTAGGTC AGCAGTAACC TTGAATTAC GTGTACTTTC TTGCATTCA	12480
CTAGCTAGCG ATAGGCAGATT TGCAACCAGTC AAGACCACTG ATACTTCTGA AGCAAAACCG	12540
CTAATAAAAT ACTTATCACT ATTATAGCGT ATGTCATAG GGACATTGT TACTGTATTA	12600
GTATACCTTT CCGTTTTAC CTGGCTAGCA CTGGTACTGT TTTGAAAATT CGTCGCCGTA	12660
GCATAGACAA ATAAGACACA AGCAAAAAAG AGTGAGGATA TGATATATAA ACTATTTTT	12720
TTCATGTTTC CATCCCTCA GCAATCGTTC TTAAAAACTA AGACCCACTT CCTCTTTGG	12780
AAGTAAGATT TCACGTAATT CTGTTCAAA TTCACTCAAGT GTTAGGTTGT GCTTAAACCT	12840
TCCATTATAG GTTATCGAAA TTCCCTCCGT TTCCCTCTGAT ACGACAAAAG TCAAGGCATC	12900
TGAGACTTCT GATAAACCAGA TAGCCGCCGG GTGTCTGGTC CCAAATTCT TGGAAATCCC	12960
TGTGTTTTT GTCAAGGGCA GATAGGCAGA CGTCACAGCG ATACGTTCTT CTTTGATAAT	13020
CACCGCACCA TCATGTAGGG GAGTGTGGG AATAAAAATG TTAATGAGAA GTTCTGCAGA	13080
AATCTTAGCA TCCAAGGGAA TTCCCTGTCGA AATATACTCC TGCAAGGTAC GTACACGCTG	13140
AATAGCAACC AAGGCCCGA TTTTACGAGG ACTCATGTAT TCAACAGACT TAACAAAGGC	13200
ACGAATCATC TGTCCTCACTAATAGG GGCATTGGAA AAGAAATCTG TCGCTCTTCC	13260
CAAACGTTCC AAACCAAGTCC GAATCTCTGG AGAGAAGATA ACAACCGCCG CAATAACCCC	13320
ATAAGTAATA ATTTGATTGA TTAACCAAGA AATCGTAGTC AAACCAATCA TATTGCAAG	13380

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GATTTGAGCT AAAATAAACCA CCAAAACTCC ACGTACCAAA ATCATAATCT TGTTCCCTGC	13440
AATAGCTTT GTAAAATGGT ATAAAATATA AGCAACAATC AAAATATCAA TCAGATTGAT	13500
AGCTATCGTC CATGGACTTG CAAACAAACT GGTCCAATAT TGCAAGATTGG ATAATTGTG	13560
AAAATTCACTC CCTGATATCC TCCCTATCAA AACACTTCG TCCTATTATA CCATTTCTG	13620
GCATTTTTT CCCTATCCTA GTCCATTATA CATTGAACAA AAATATGATA AAATAAACTG	13680
ACTAAAAAAA ACAAAGGAGA AACTATGTCT CAACTCTATG ATATTACCAT TGTGGGTGGT	13740
GGTCTGTGCG GGCTTTTGC AGCCTTTAT GCCCACCTAC GCCAAGCCAA GGTTCAAATC	13800
ATCGACTCTC TTCCCCAGCT AGGTGGACAA CCTGCTATTG TCTACCCCTGA AAAGGAAATC	13860
CTAGACGTAC CAGGCTTCCC AAACCTGACT GGAGAAGAGT TGACTAACCC CTTGATTGAA	13920
CAGCTAAATG GATTGATAC CCCTATTGAT CTCATGAAA CGGTTCTTGA GATTGACAAA	13980
CAAGAAGAAT TTGCCATCAG AACTTCTAAA GGAAGTCACC TGACTAAAC AGTTATCATC	14040
GCTATGGGTG CGGGTGCCTT CAAACCACGT CCGCTGGAAC TTGAAGGGGT TGAGGGCTAT	14100
GAAAATATCC ACTACCACGT TTCTAACATT CAGCAATACG CTGGTAAGAA AGTGACGATT	14160
CTTGGTGGGG GAGACTCGGC TGTGGATTGG GCTTTGGCTT TTGAAAAAAAT CGCACCAACT	14220
ACCCCTGTTG ACCCGAGAGA TAATTTCCGT GCCTTGGAAC ACAGTGTCA AGCCTTGCAA	14280
GAATCATCTG TAACCCTCAA GACACCATTG GCCCCTAGCC AACTCCTTGG AAATGGAAA	14340
ACACTTGATA AACTTGAAAT CACAAAAGTC AAATCTGATG AAACCTGAAAC CATTGACCTA	14400
GACCACCTCT TTGTCAACTA TGGTTTCAA TCTTCTGTCG GTAAACCTTAA AAACCTGGGG	14460
CTCGACCTCA ACCGTACAAA GATTATCGTC AACAGCAGAAC AGGAATCCAG CCAAGCAGGT	14520
ATCTATGCTA TCGGTGACTG CTGCTACTAT GACGGAAAAA TTGATCTGAT TGGACAGGC	14580
CTCGGAGAG CTCCAACCTGC TGTCAACAAAC GCTATCAACT ACATTGACCC TGAACAAAAA	14640
GTACAACCAA AACACTCTAC TAGTTTATAA AAAAGAACCA CGAGTCACAT AGGATTCTG	14700
GTTTTATAAT TCATCCGCTA TCTTATTGAT TTTTCTGAGT CTGTGATTGA CACCACTTTT	14760
GGTCAGAGGG GTGCTGAGAC TATCTGCTAA CTGCTGGATA GAGTAGTCTG GGTGCTGAAT	14820
CCTCAGTTGC GCCACTTCCT GCAAATCTAC TGGCAAAATT TCTAAGCCCA TGATATCTTT	14880
GATTTTACTG ATATTGTTAA TGGTCTTCAT GCTGGCAGAA ACTGTCCGAG CGATATTAGC	14940
TGTCTCGGCA TTATTAGGCC GATTGAGGTC GTTACGGGTT TCTCGAAAAA TCTTAACCCG	15000
CTCAAAATCA TCACGTGCCT GCATGGCTCC TATTACTATC AAGAAGTCCA TAATGTCTTC	15060
TGCTCGCTGG AGATAGGTCA CAGCCCCCTT CTTGCGCTCA AGCACCTTGG CATCCAGTAA	15120
AAACTGTTGG AGAAGGGAGG CAATTCCCTG CGCGTGGTCC AGATAAACAG AACTGATTTC	15180

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CAAATGGTAC TTGCGCTGACT CAGGGTCACG AATGCTCCCA TTTGCCAAGA AAGGCCACA	15240
GAGATAGGCA CGACCTGCTT CCTCATCCGA TAAAATCGCC TCATCAATAC CTGTTTCCAG	15300
GCCAAAGAAA GAGTCTGCCA AGTGCAAATC ACTTAACAAA TCCTGCACCT TTTCATCTGT	15360
AAAAACGGTA TAGACCGCGAT TCTTGCAGG ATTGCTCCGT TGGTGGTGAC GAATTCAGA	15420
TTTGATTTCA TAGAGATGGA GAAAGGACTC ATAGAGGTGA CGGGCCAGTT TGGCATTTTC	15480
TGTCACAACT GACAAAGTCA AGCCCGAAGT CGAGAGACCG ATGCTACCAAG ACATTTTGAT	15540
AATGGCAGAT AATTCACTGCC AGCTCAGATG GTGTTGGCCC AGGATTTCTT CTTTTACTGC	15600
TACTGTGAAA CTCATTTTT CACCTGTATA ATGCCGATCA ACTCGTCCAC AATCAAATCT	15660
CCATCGTGGAGGACCCGCCTTCCAGA CGAAGGAAGT TAGATGAAAT CACGCCGCAA	15720
ACTTGCTTAC AAAGACCTAC AAAATCGTGT TCCACTTGCA CTAAGTATTC ATCAAAACGG	15780
TTGGAATTCA TGTATTCCCTG AGGCACTTTT TCAATATTCA CCAAGACAGT GTCGATAAAA	15840
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TCCCCACGTT GGGTCATGAT ATTGCGAGACA TAGGCAATTCTGCTT TTCCAAAAGA	15960
GCCCCCCCAA TTTCCTTAAT CACGATATTG GGCAAAATAG AGGTAAAGAG GGAACCTGGC	16020
CCTAGGACAA TCATGTCACT TTCAAGGATG GTCTGCACTA CTCGACGGCT GGCCAGAGGC	16080
GTATCATCGT TTAGGGCATT GGTACACATAG ACATTGTCAA TTATGCCTCG ATGGTCTACA	16140
ATATGACTCTCTCAGCCAC TTCTGTCCCA TCCCTGAAAGA CTGCATGAAG GGTCAAAGGA	16200
TGGTCACTGG AAGGATAAAAT TTCCCTGTGT GTATGGAAAA ATTTGCTCAA TAACTGCATG	16260
GCATTATAGG TTGAAACCTTG CATTCTGAC AAGCCAGCAA TGATGAGATT TCCCAATGGA	16320
TGGCCAGCAA AGGCTCCGGC ATCCTCAGAG AACCGATACT GAAAGACCTT CTCATAAAC	16380
TTAGGCATAT CCGACATGGC CACAAGGACA TTACGAAGAT CACCTGGCGG TGTCAACTGT	16440
TGCATATTTT TTGGAGTTC ACCTGAAGAA CCACCATCAT CTGCCACCGT CACGATAGCT	16500
GCGATTCCA CATCTTTTC CCGCAGACTT TTAGAATGA CGGGACTTCC AGTCCCTCCA	16560
CCAATCACCG TTATCTTGG TTCTCTCATG AACGGTTAC CGTTTCCCTT CTGCGGTCTT	16620
TGTCGGGATG CCCTTCATTA ACAGACCAAT TCTTGGATAA GTCTGCGGCC AAGCGTTTAG	16680
CAAATGCCAC ACTACGGTGT TGTCCACCCG TACATCCCAT GGCAATGGTC AAAACGGACT	16740
TACCTTCCTT TTGGTAACCTT GGCGAGATCG GCTCAATCAA GGCAATAAAA TGTTGATAAA	16800
AGTCTTCTGA CTCAGGATGG TTCATGACAT AATCATAAAC AGGTTCATCC ACACCCGTTT	16860
GGTTTCTCAG TTCTGGTAAA TAATAGGGAT TTGGCAAGAA ACGGACATCA AAGACCAAGT	16920

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CCGCATCAAT CGGGATTCCA TACTTAAATC CGAAAGACAT GACTTCGATA CGGAAAGACT	16980
GGGCTTGTTC TTGGTCTGAA AACTGCTCTG CAAGGGTTTT GCGCAGCTCA CGTGGAGTGA	17040
GTTCACTCGT ATCCACCACA TTTTGGCTCA TATTTTCAA AGGTGCCAAG AGTTCACGTT	17100
CCAACTTGAT TCCATCTAAA ATACGACCGT CTGCTGCTAG TGGGTGACTC CGTCTGGTTT	17160
CCTTGTAAACG ACCGACCAAT TCCTTATCAG CCGCATCCAA AAAGAGGATT TTGAAATCCA	17220
AACCATCTTG ATTTTCCAAC TCATCCAAA CAGCTTGAAT CTCTGAAAAG AAAGAACGGC	17280
TACGCATATC CACTACCAAG GCCAACTTAG GATTGTCTTC CTAAATTCA ACCAGCTGCA	17340
AAAACCTTAGG CAAGAGAGCT GGCGGCATAT TATCAATGGT GAAATAACCT AGATCCTCGA	17400
AGGACTGAAT GGCTACAGTT TTCCCTGCGC CACTCATCCC TGTCACAATC ACCAAAGTGA	17460
GTTGTTCTT TGTCATCTT TTCTCCTTAT ATCAAAAGAA GTTGGCAAC ACCAAACTTC	17520
AACTAGCTTA TCCAATCTCT GCGATGACTT CAATTTCGAC TTTTACATCA CGAGGAAGAC	17580
GAGCTACCTC CACAGCTGAA CGAGCTGGGA ATTCCCTTT GAAGGCCGTT TGGTAAACCT	17640
CATTAAAAGG AACAAAGTCG TTCATATCGC TCAAGAAGCA AGTTGTTTG ACAACATGGT	17700
CAAAGTCTGT TCCCTGCTCT GCCAAATAG CACCGATGTT TTTCAAGACT TGCTCTGTCT	17760
GTTCTGGAT ATTCTCTCCT ACAATTTCCTT CAGTTTCAGG GGATAGGGGA ACTTGACCGC	17820
TAGCAAACAA AAGGTTGCCA ACGATTTTC CTTGAACATA GGGTCCGATA GCCTTGGGG	17880
CCTTATCTGT ATGAATTGTT TTTGCCATT TCTTTTCCTC ACAATTTCCT TAAGATTGCA	17940
TCCCAAGCCT CATCCATCCC TGCCCTACTG ACAGATGAAA AGAGGATGAA ATCGTCACTC	18000
GGGTCAAAGT TTAATTCTT TTTGATTGCT GATTCTGCT TGTTCCATT ACCACGAGGA	18060
ATCTTGTCGG CCTTGGTCGC CACAATGATG ACTGGAATCT CATAATACTT GAGAAATTG	18120
TACATCTGCA CATCATCTGC TGACGGGTCA TGACGAAGGT CAACTAGACT GACAACCGCA	18180
CGGAGATTT CCCGAGTCGT TAAGTACTCC TCAATCATGC ACCCCCCACTT TTCACGTTCC	18240
TTTTTAGAAA CACGAGCATA GCCATAACCA GGCACATCCA CAAAGCGCAT CTGTGTCATCA	18300
ATGTTAAAAA AGTCAGGAG CTGGTTTTA CCAGGTTTC CTGATGTACG GGCGAGATT	18360
TTACGGTTCA ACATAGTGT GATAAAGCTG GATTACCAA CATTGAAACG CCCTGCTAGG	18420
GCAATCTCTG GCAGTTCATC CTGGGGATAG TGGGACTTAT TAGCTGCACT GAGCAAGATT	18480
TCAGCATTGT GTGTATTAAG TTCCATAGTC ACCTCTAGGC TGTTTCTAGG ATCGGTTAT	18540
CCGTTCCATC TACAGTTCT TTAGTGATGC GAACCAATT CACATTTC TGACTCGGCA	18600
CCTCAAACAT GACATCTAGC ATGGTTCTT CGATGATGGA GCGAAGTCCA CGCGCCCCCTG	18660
TCTTCGTTTC GATTGCTTTA TTAGCAATCT CTTGAAGGGC TTGCTGTC AATTCCAAC	18720

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CAACATCATC ATAAGAAAAG CAGGTTTGGT ATTGTTTCAC CAAGGCATTT CTTGGCTCTT	18780
TCAAGATGCG AACCAAAGTCA TCAACGGTCA ATTGCTCAAG AGCCGAAAA ACAGGCAAGC	18840
GTCCAATCAA CTCAGGGATA ATACCAAATT TTGGAATGTC TTCAGCGATG ATTTCTTGCA	18900
TGTATGAGCT GTTTTCGTCA ATGCCATTAT TATTTGACC AAATCCGATG ACTTTTCAC	18960
CCAGACGTTG TTGACAATT TCTCAATAC CATCAAAAGC ACCACCCCAGC ATGAAGAGGA	19020
TATTTTTGT ATCCACTTGA ATCATCTCTT GTTGTGGATG TTTGCGTCCA CCTTGAGGCG	19080
GTACGCTAGC AACAGTTCCC TCAATAATCT TGAGAAGGGC TTGTTGCACC CCTTCACCAG	19140
AAACATCACG TGTGATAGAC ACATTCTCAC TCTTCTTGGC AATCTTGTCA ATTTCATCCA	19200
CATAGATAAT GCCACGCTCT GCACGTTCGA TGTAAAGTC AGCAACCTGC AAGAGTTTGA	19260
GGAGGATATT TTCCACATCC TCACCCACAT AACCAGCCTC CGTCAGAGCT GTCCGATCCG	19320
CAATAGCAAA AGGTACATTC AAGCTCTTAG CCAAGGCTG GGCAAGGAAA GTTTTCCCTG	19380
AACCAGTTGG GCAATCATC AAAATGTTG ACTTCTGCAA ATCCACATCT TCTGACTCTT	19440
CGCGTGTATC GTGAAATTG ATGCGTTGT AGTGGTTATA AACCGCCACT GCCAAGGCAC	19500
GCTTGGCACG ATCTTGACCA ATTACATAGT GGTTCAAGAT ATGGAGGAGT TCAATTGGTT	19560
TTGGCACCTC AGACAAGTCT GCCAAGACTT CCTCAACCAA TTCTTCTCGA ATGATTTCCCT	19620
GAGCTAACTC CACGCATTCA TTACAAATAA AAGCATTGTT GCCAGCAATT ATTTTTGTA	19680
CTTCTTCTTG GTTTTGCCA CAAAATGAGC AATAAACCAT CATATCATT TTTCTATTTG	19740
TAGACATGAT TTCCCTCCAT TCTATACTGT CATTCTATCT AAAATAAGGT CATGAAAAAA	19800
GCATGAATAC TATTGACCAAG ATTGGTAAAG GCATTTAACC AAAGGAGGAT AGAAAGCCCG	19860
TAACGCTTTT TACGAAAGC TTGTGCTCCT GCCAGAAAGC AGATGAAACA CAGAAAAGCC	19920
GTGAATAGAC CAAATAAACT CCGTCCATT AGACTTCCTT TCTCTGCGG TATTGGATGG	19980
TAAAATCATA AGGATTCTTC TCATCTTGG CGTAAATTT GCTTGAAACT GTCTAAAAAA	20040
GAGACAAGTC AAGTTCTCA GGGAAATAGG TATCTCCCTC CACCCGAGCA TGAATGTGAG	20100
TGACAATCAC TTCATCAAGG TAAGGTTCAA AAGCCTGAAA AATTTGCTTC CCACCGATAAA	20160
TGTAGAGATT CTTTCTTGA GCCTGATACC AGTCAAGAAC AGACTGGACG TCCTGAAAAG	20220
TAGCAACCCC ATCTATCTTT TCTTCCGGAT TACGCGTCAA AATCAAGGTT TCCCGTTTG	20280
GAAGCAAGCG ACGCCCCATC CCATCAAAGG TCACACGCC CATCAAGATA GCATGATTCA	20340
GAGTTGTTTC TTTAAAGTGC TGCAATTCTG CTGGCAAATG CCAAGGCAGA CGATTTCCCT	20400
TACCAATCAC ACCCTCTTCA TCTTGGGCC AAATAGCTAC GATTTCTTA GTCATGCTTC	20460

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CATCCTTTTC ACTGATAGTA CTATTTTATC AAAAAACTCA AAAAAAGACT GGTTTGGAAAT	20520
AGCTTACAAA ATAGAAAAAA TCTGTAAGAA ATTTCCCTACA GATTTATCTA TGTTTCCCTTA	20580
TTTCTTACAA ACCAGGTGCT TGCCAAGTT CGGCTGCAAG CATCCAAATT GTTTTATCTG	20640
TTTCAGTTTT AGCCGCTGCA AAGATAACCGT TTGTCACATC GTCACCTCT TCATCAGTGA	20700
CATCCAAACC TTTTGGAAA AGTTCTGACA AGTAACGGTA GATAACAAGA ACACGTTCCA	20760
AGCTTCTTC AACATTACGG TATTCACCAG CTTCTTCTTC GATTTCACTA TTTTGAAGGA	20820
ACTCTGTCAA TGTAGAGAAT GGGCTTCCAC CGAGTGTAAT CAAGCGTTCA CTGATTTCAT	20880
CCAATTGACC GTCAAGAGCT TCCATGTACT CATCCATTTC TGGATGCCAT ACAAGGAAAC	20940
CACGACCATG CATATACCAG TGCACTTGGT GCAAAGCAAC GTGAGCTACA TACAAATCAG	21000
CAACAGCTTG GTTCAAGACT TCCCTTGTAA TTGCCAATGC	21040

(2) INFORMATION FOR SEQ ID NO: 56:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 2387 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 56:

ATTCTTAATA CGATTAAG GCTTATTACT AAAAGAAAAT TTCAGTTAGA TGAACCAAAC	60
TTGCTCGTCA AATCCCGATT TAACGAGATG TTTGGGGAAA ATAAAATATT TGAAAGCATT	120
GATAACTTAT TTGATATTAT AGATGGTGT AGGGGCAAAA ATTATCCTAA ATCAGATGAG	180
TTGTTTAGTG AGGAGTACTG TTTATTTTA AATACAAAAGA ATGTTACTAA AAACGGATT	240
TCATTCGATA CAAAGCAATT TATCACTAAA ACAAAGGATA AATTACTTCG AAAAGGCAA	300
CTTGAGCGTT ATGATATACT CTTGACAACA AGAGGTACTG TTGAAATGT AGCGTACTAC	360
GATGAATTAA TAAAATATAA ACATTTACGT ATAAATTCA GTATGGTAAT ATTACGTCCC	420
AAGACACCAA ATCTAAATCA GAAATTATT ATCCATGTAA TAAGGAATAA TAATTATAGT	480
CGAGTGATAT CAGGAAGTGC TCAGCCTCAG TTACCAATTAA CAAAATTAAA AAAAATACTT	540
CTCCCCCTCC CCCCACTAGC CCTCCAAAAT GAGTCGCAG ACTTTGTAGT CCAGGTCGAC	600
AAATCACAAAT TGGCAATCCA AAAATCTCTG GAAGAACCTTG AAACTTGAA GAAATCTCTG	660
ATGCAGGAGT ATTTTGGCTG ATATTCTGCC ATTGTAATTAA CGGTAATGAT TTGTTATAAT	720
ACTTCAAAGG AGGAAATCAG ATGGTAGTAA AAACAAGAAA ACAAGGAAAT TCAATCACCA	780
TTACGATTCC AAGTGAATTAA ATATTCCTAA GTGGTGTAA ATACGAAGCG AAATTGTTAC	840

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CAAGTGGTGA GATTATCTTT ACTCCTGAAG AATTGGGCA GCAGGTTCT TATGTATCTG	900
ATGATGCCCTT TGACTTAAAT TTAGATAAAA TATTTGACGA ATACGACGAT GTTTCAAAG	960
CTTTGGTGG AAAATGACAA TCTATTGAC AGAAAAGCAA ATTGAAAAAA TAAATGCTTT	1020
AGCAATTCAA CGGTATTCTC CAAATGAGAA AATTCAAACA GTTAGTCCTT CTGCCTTAAA	1080
TATGATTGTG AACTTACCAAG AACAAATTGTG CTTTGGGAAG CCTCTTTATC CAACAATT	1140
TGATAAAGCA ACGATACTAT TTGTCATT GATAAAGAAG CATGTTTTG CTAATGCTAA	1200
TAAAAGAACT GCTTCTTCG TTTTGGTCAA ATTTTACAA TTAAACGGCT ATCGTTTTTC	1260
TGTAACCGTA GAAGAACAG TAAAAATGTG TCTAACCATC GCAGTAGAAG CTTAACTGA	1320
TGAAAAATG ACAAGCTACT CCAAATGGAT TTCTGAACAT TCTGTTAGAG AAAAGGTCAA	1380
AAAGTAACCT AGTATGCTGG ATTTGAATGA GCACAAGAAA ATAATGAAC AGACAATATT	1440
AGAATTCTGT AATGCAGAAA CTGATATTGT CTCTTTTAT TGATGAATAA GAAAGTGAGA	1500
AATTATGGAA TCAAAAGTTA CAATTATCAT GCAAGAAATG TTACCTCTTT TAAATAATGA	1560
ACAATTACTA GCGTTGAGAG AGAGTTAGA ACATCATCTA GTAGACGGAA AAAAGCAGCA	1620
GAAGTATTG AATAATAACC TGTGCAACT ATTTATTACC GCCAAGCAGG TAGAGGGCTG	1680
TAGCTCAAAA ACAATTGCTT ATTATCAGAG GACGATTGAA AACTTGTAA ATGCTATTAA	1740
AGAGTCTGTG ACACAACCTCA CAACAGATGA TTTAAGGAGT TATTTAGCAA ATTACCACTC	1800
TGAAAAGGAT TGTAGTAAGG CAAATTAGA CAATATTAGG CGTATATTGT CTTCTTTTT	1860
TGCTTGGCTT GAGCAAGAGG ATATATCATT AAAATTCCC TTGACGGAT ACAGAAAATT	1920
AAGACTGAGC AAAATGTGAA GGAAACCTTAT ACTGATGAAC ATTTGGAAAT TATGCGTGAT	1980
AACTGTGAAA ATTTGAGAGA TTTGGCAATA ATAGACCTAC TAGCATCGAC AGGTATGCGT	2040
GTAGGGGAGC TTGTACAGTT GAATCGTTCA GATATTGATT TTGAAAACAG AGAGTGTGTT	2100
GTCTTGGTA AAGGAAAGAA GGAGAGACCA GTATATTGT ACGCTCGTAC GAAAATTCA	2160
TTAAGAAATT ATCTTAACGA CAGAAAAGAT AGTCACCCCTG CTCTTTTTGT AACGCTAGTT	2220
GGAAAAGTCC AGAGGCTTGG AATGCTGGT GTAGAGATTC GCTTAAGAAA GTAGGAGAC	2280
AAACTCGCCA TACAAAAGGT TCACCCACAT AAGTCAGAA GAACTTAGC GACTAAGGCA	2340
ATTGATAAAG GTATGCCTAT CGAACAAAGTC CAAAAACTGC TAGGTCA	2387

(2) INFORMATION FOR SEQ ID NO: 57:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 10669 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double

504

(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 57:

ATATTAAAGC GACTTTCTGT GCGCTAGGGA AAAATGTTCC TGGGAATGAG GACTTGGTGA	60
AGAGGATAAA ATCTGAAGGT CATGTTGTTG GAAACCATAG CTGGAGCCAT CCGATTCTCT	120
CGCAACTCTC TCTTGATGAA GCTAAAAAGC AGATTACTGA TACTGAGGAT GTGCTAACTA	180
AAAGTGCTGGG TTCTAGTTCT AAACTCATGC GTCCACCTTA TGGTGCTATT ACAGATGATA	240
TTCGCAATAG CTTGGATTTG AGCTTTATCA TGTGGGATGT GGATAGTCTG GACTGGAAGA	300
GTAAAAATGA AGCATCTATT TTGACAGAAA TTCAGTATCA AGTAGCTAAT GGCTCTATCG	360
TTTGATGCA TGATATTCA C AGTCCGACAG TCAATGCCCTT GCCAAGGGTC ATTGAGTATT	420
TGAAAAATCA AGGTTATACC TTTGTGACCA TACCAAGAGAT GCTCAAACT CGCCTAAAAG	480
CTCATGAGCT GTACTATAGT CGTGATGAAT AAGCAAGAAA AAATAGGTCT GTTAGATATT	540
TGACAGACTT ATTTTTACA GAATATAGTA CTACTAAAAA AATGTTTAT GCTATAATTG	600
ATGAATAAAA TAGAAGGAGA AGCATATGAA TACCTATCAA TAAATAATG GAGTAGAAAT	660
TCCAGTATTG GGATTGGAA CTTTTAACCG TAAGGATGGA GAAGAAGCCT ATCGTGCAGT	720
GTTAGAAGCC TTGAAGGCTG GTTATCGTCA TATTGATACG CGGGCGATTT ATCAGAATGA	780
AGAAAGTGTG GGTCAAGCAA TCAAAGATAG CGGAGTTCCA CGTGAAGAAA TGTTCGTAAC	840
TACCAAGCTT TGGAAATAGTC AGCAAAACCTA TGAGCAAAC CTGCAAGCTT TGGAAAATC	900
TATAGAAAAA CTGGGCTTGG ATTATTTGGA TTGTGATTTG ATTCAATTGGC CGAACCCAAA	960
ACCGCTCAGA GAAAATGACG CATGGAAAAC TCGCAATGCG GAAGTTTGGG GAGCGATGGA	1020
AGACCTCTAT CAAGAAGGGG AAATCCGTGC TATCGCGTT AGCAATTTC TTCCCCATCA	1080
TTTGATGCC TTGCTGAAA CTGCAACTAT CGTTCTGCG GTCAATCAAG TTCGCTTGGC	1140
GCCAGGTGTG TATCAAGATC AAGTCGTAGC TTACTGTCGT GAAAAGGGAA TTTTATTGGG	1200
AGCTTGGGG CCTTTTGGAC AAGGAGAACT GTTTGATACG AAGCAAGTCC AAGAAATAGC	1260
AGCAAATCAC GGAAAATCGG TTGCTCAGAT AGCCTTGGCC TGGAGCTTGG CAGAAGGATT	1320
TTTACCACTT CCAAAATCTG TCACAAACCTC TCGTATTCAA GCTAACTTGT ATTGCTTGG	1380
AATTGAACTG AGTCATGAGG AGAGAGAAAC CTTAAAAACG ATTGCTGTTC AATCGGGTGC	1440
TCCACGAGTT GATGATGTGG ATTCTAGAA AATCATAAAA AGAATTGTAC ATTATTCTAA	1500
TTTTGATAT AATAGTCAGC AGGAAAGAAA GTCTTATGGC GTTCTTCAAG CGAGCTTGGG	1560
ATAGTGGGAG CCAAGTAGGG CAAAATAAG GGCTGGCGCT TTCTGTAGTA TTTTCAAAAA	1620

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CAATGAAGTA ATAAATTAGG GTGGAACCGC GTTTCTGACG CCCCTAGGTT AAATCAACCT	1680
AGGATTGTCA GATGTGGTTC TTTTGCTTAT TCAGTCTATT GTGTGAAAGA AAGGAGAGCC	1740
GTGGACAACC TTTATCTTGT AAAAGACGAT AGTCAACTAG CTACATTTCG TGATTTGTA	1800
GTAAGAAAATA CTGAAAAGTT GAAAGATTAT CAATCTTTT TAAAGAATGA ACTTGCAGTC	1860
TGTGATTAC CGCAAGCTGT TATTGCTCA GATTTAATG CTGCTACACA GATTATTAGG	1920
GAAAGTGCTG TTCCAACCTA TACAAATAAT AGACGAGTGG TTATGACGCC TGATTTAGCT	1980
GTTTGGAAAG AATTGTTATT GTATCAGTTG ATGGACTACG AGTGTCTGA GCAAACTCAA	2040
GCAATAGAAA GTCACTATCA TTCTTTATCT GAAAATTTC CTTTACAGAT TGTAGGACAT	2100
GAGTTAGCTC ATTGGTCGGA CATTTTTAG ATGATTTGA TGGTTATGAC TCTTATATCT	2160
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TTCAAGCGGA AAAAATTGTT AATCAATCTC TCGTAGAACT TTTTCAGAAC AGATATAGTT	2280
GGCATTCATT GAATGATTTT GGTTCTTCGA CTTATGATAA GAACTATGCA AGTATTTTT	2340
ATGAATACTG GCGCAGCTTT TTGACAGTAG ATAAGTTGGT AGAAAATTAA GGTAGTGTAC	2400
AAGCGGTCTT AGATTCTTAT CATTATGGG CAAATACAGA AAAAACTTTT CCCTTGTTAG	2460
ATTGGTTGT TCAGCAGAAA TTAAATTGAA AAGAAATATA AAAACTAAAG GAGTAAACAA	2520
TGTCTAAGAA ATTAACATTT CACTGCATCA GTGGCAGAGA CCTCCTTACA GTCGGGCTGC	2580
TCCACGCTCA GCACTAGAGT GCCTGAGCTA GACCGAGTAC TAACTCGTCT TGCCCTCGTAT	2640
GATCGACGAG GCAGACTCGT GTCGCAAGTA ATTATTTTTT ATTAAGGAGT ATTCAATGTC	2700
TAAGAAATTAA ACATTCACT GCGTCAGTGG CAGAAACCTC CTTACAGTCG GACTGCCCTA	2760
CGCTCAGCAC TAGAGTGCCT GAGCTAGACG CAGTACTAAC TCGTCTGCC TCGTATAATC	2820
GACGAGGCAG ACTCGTGTGCA CAAGAAATTAA TTTTTTATTA AGGAGTATTC AATGTCTAAG	2880
AAATTAACAT TTCAAGAAAT TATTTTGACT TTGCAACAAT TTTGGAATGA CCAAGATTGT	2940
ATGCTTATGC AGGCTTATGA TAATGAAAAA GGTGCGGGGA CAATGAGTCC TTACACTTTC	3000
CTTCGTGCTA TCGGACCTGA GCCATGGAAT GCAGCTTATG TAGAGCCATC ACGTCGTCT	3060
GCTGACGGTC GTTATGGGA AAACCCCTAAC CGTCTCTACC AACACCCACCA ATTCCAGGTG	3120
GTCATGAAGC CTTCTCCATC AAATATCCAA GAACTTTACC TTGAGTCTTT GGAAAATTG	3180
GGAATCAATC CTTTGGAGCA CGATATTGCGT TTTGTTGAGG ACAACTGGGA AAACCCATCA	3240
ACTGGTTCACT CGGGTCTGG TTGGGAAGTT TGGCTTGACG GAATGGAAAT CACTCAGTTC	3300
ACTTATTTCC AACAAAGTCGG TGGAATTGGCA ACTGGCCCTG TGAUTGCGGA AGTTACCTAT	3360

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GCTGATGGTG TAAAATACGG AGAAATCTTT ATCCAGCTG AGTATGAGCA CTCAAAATAT	3480
TCATTTGAAA TTTCGGACCA AGAAATGTTG CTTGAAAACT TTGATAAGTT TGAAAAAGAA	3540
GCTGGTCTG CATTAGAAGA AGGCTTGGTA CACCCTGCCT ATGACTATGT TCTCAAATGT	3600
TCACATACCT TTAATCTGCT TGACGCGCGT GGTGCCGTAT CTGTAACAGA GCGTGCAGGC	3660
TATATCGCTC GTATCCGTAA CTTGGCCCGT GTCGTAGCCA AAACCTTTGT CGCAGAACGC	3720
AAACGCCCTAG GCTACCCACT TTTGGATGAA GAAACAAGAG CTAAACTCCT AGCAGAACAC	3780
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TGCCGCAGTG ATGAAGGTAT CCTTAGTGAA ACTAAGGATA CTAGGCAAAA TTGGAGACTT	3900
TTGGCTCCAA TTTAGCAAT GAAACAACGA AGTTGGTTCG TTGCGTGCCA ATCACATAAG	3960
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CGAAAAAAATG GCAGCCTTCC TCAAGGGAAA ACCCCTGTCT TTTGAAGCCA TTCAAACTTT	4140
CTCAACACCA CGTCGTTGG CTGTCGTGT AACTGGTCTT GCAGACAAAC AGTCTGATT	4200
AACAGAAAGAT TTCAAGGGTC CAGCAAAGAA AATTGCCTTA GATAGTGATG GAAACTTCAC	4260
CAAAGCAGCT CAAGGATTG TCCGTGGAA AGGTTTGACT GTTGAAGATA TCGAATTCCG	4320
TGAAATCAAG GGTGAAGAAAT ATGTCTATGT CACTAAGGAA GAAATTGGTC AAGCAGTTGA	4380
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TGAGCAAGAG TTTGACTTGG ATTCCTTGA TATCAAGGGAA AGTCGTGTGA GTCGTGGCCA	4560
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TAAGCAGTTT GTAATCGCAG ATCCATGTGA ACGTGAGCAA ATGATTGTTG ACCAAATCAA	4680
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TGATCAAGAT GGAAACTCT TGCCAAACTT CATTCTGTT CGTAACGGAA ACGCAGAGCC	4920
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ATTCTCTGG CGTGAAGACC AAAAATTGGT GATTCAGAT CTTGTTGAAA ATTAAACCAA	5040
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CACTGTACTT TTGGCAGAAA AAGCTAGTTT GTCAGTGGAT GAAACAGTTG ACCTTGCTCG	5160

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TGCAGCAGCC ATTTACAAGT TTGACTTGTT GACAGGTATG GTTGGTGAAT TTGACGA	5220
CCAAGGAATT ATGGGTGAAA AATACACCCCT TCTTGCTGGT GAAACTCCAG CGGTGGCAGC	5280
TGCTATTTCGT GAACACTACA TGCCCTACATC AGCTGAAGGA GAACCTCCAG AGAGCAAGGT	5340
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ATTGATTCCA TCAGGTTCTA ATGACCCCTTA TGCCCTTCGT CGTGCACACTC AAGGTGTGGT	5460
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TGCATTGAAA TTTGACAGTT TGACTTATGA AAATAAAAGCA GAGGTTATGG ACTTTATCAA	5580
GGCTCCGTGTT GATAAGATGA TGGGCTCTAC TCCAAAAGAT ATCAAGGAAG CAGTTCTTGC	5640
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CAAGGAAGAA GATTTAACAC CATCTGTTGA ATCACTTTCT CGTGCCTTTA ACCTGGCCGA	5760
GAAGGCAGAA GGTTTGCTA CGGTTGATTC AGCACTATTT GAGAATGACC AAGAAAAAGC	5820
TTTGGCAGAA GCAGTAGAAA CACTCATTGATCAGGACCT GCAAGTCAGC AATTGAAACA	5880
ACTTTTGCG CTTAGCCCCAG TCATTGATGC TTTCCTTGAA AATACTATGG TAATGGCTGA	5940
AGATCAGGCT GTCCGTCAAA ATCGTTGGC AATCTTGTCA CAACTAACCA AGAAAGCAGC	6000
TAAGTTTGCT TGTGTTAACAC AAATTAACAC TAAATAAAAT TTGATAAACG GACTTTATCT	6060
TATTACAAAG GAGAAGAAAT GGATCCGAAA AAAATTCCCTC CTATCAATGA GCTTGCTAAA	6120
AAGAAAAAAA CAGAAGGCTT AACACCAGAA GAAAAAGTGG AACAAAGCCAA ACTACGTGAG	6180
GAGTACATCG AAGGTTATCG CGCGCTGTT CGTCACCACA TTGAAGGAAT CAAAATTGTG	6240
GACGAAGAAG GAAACGATGT TACACCAGAA AAACATACGCC AAGTACAACG TGAAAAAGGA	6300
TTACATGGCC GTAGTCTTGA TGATCCAAT TCATAATAAT ACTCTTCGAA AATCAAATTC	6360
AAACCAAGTC AGCTTCACCT TGCCGCTACTT AAGTACAGCC TGCGGCTAGC TTCCCTAGTT	6420
GCTCTTGAT TTTCATTGAG TATATGTATT CTTCTTTTA ACAAAAGATAG ATGAAACGAT	6480
AACAAAGAGA CTAGCAGTT GTGTTGCTA GTCTTTTTC GCTAAAAAAG GAACCATAAT	6540
GGTTCTAAA AACTATCATT AGTAACCTGC ACCGGCTGTA CGCTCTCGT CACCACCGTG	6600
GCCTCCAGCA TCCCCTGAAT CAGAAGCGCC AGAAGTAGCA TCGCGCTCTC CATGACCTCC	6660
GGCAGCAGGA GCAAATGGTC CGCTACCACC CACCAAACGT TGACCAGTCT CTTTTAGGTA	6720
CCAGTCAAGC CATGGTTGGA AGTTAAAGAC GATTTCATTG ATACCAGCGT ATGATCCATC	6780
AGGATAGTAC ATTGCTTGTT AGTTGTGAGT GTTGATAACA CCTGCAGGAG AACCTGGAAC	6840
GATCGTACGG ACGTATTCTT GGTTCCGTT GCGAAGTGTGTT CCGATAACCC ACTCTACGTT	6900

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CTTCATACGT GCTGGTGGAA GAGAACCATG AACAGTCGAC ATACGGCTAC CTGATTGAGG	6960
TGGTACACGT TTAGCGAACAA TAGTGTCTGG ATCTTGGTGA CGCTTGGTTGT AGTAGAGGAA	7020
TTGGTGTGTT CGCTCACCGT ATGTCATTC AAATGGCATA GCTTTCAAGA ACATATCAA	7080
TTGGTTAACT GTTAGGATAC CGTGGTCCAA TTTGACATAG GTATCACCAG AAACAGCACC	7140
AGTGAATGCT GCAACTTTT CTACCCATTC TGGATCGTCA GGGTCAACTT CTGTGATGGT	7200
TGTAGCGATT GGTTTCCAC AATCCAAGTC TTCTGATTG ATTGGTTTG GTTTTTCAA	7260
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AGTGCCTTCG TTGGTGATAT TTCCGTTGTT ATCAAAGCT TCCTTAGCTT TACCAAGAAG	7380
GAATTCGTTA CCTGGAAGCG TGTAGGCATT AACACCTGGA GCATCAAGGA TTTTACGAAG	7440
GTGAACCTGA GCACGTGATG TTCCCTGGTC ATAGTATGAT GCACCCACAA TCATAACAGG	7500
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TGAGATAGTG TGGTTATGCT CAGGAGTAGC AATGATAACA CCATCTGCAC GAGTAATTTT	7620
GTTATATAAA TAACGTAATT GGAAACTTTC ATCCCATTTC TCATCTTGGT TAAACATTGG	7680
AACTTCGTCA ATTTCAAGAA CTTCTAATTC AAATTGAGT TTGAAGTAGC GACGGATAAA	7740
TTCCAAGAGC TTACGGTTAT ATGATTGATC GTAGTTGAT CCAACAAAGTC CAACAAATT	7800
CATTCTTTT GGTCTCCTAT CTTACAAATT TTCCCAGTCA AAGTCTTCAG CATCTTGCG	7860
AAGTAATTCT TGTGCATTAC GTAATTTC TGTGATTTT ACAAAAGATAC GGAAGTCATC	7920
AAAGATGGCA TCCAATTCT TGATAACATC AAGGTCAACC AAGTCGCCAC TTGGGTTAAA	7980
TGCTTGAGA GAGTGTGAGA GCAAGAACCC ATCTGGAAGA ACATTTGCCT TGATTTCAAG	8040
AGCATTCAAG ATTTGACGAA GTTGCAATTG GGCACGAGAT GAACCAAGCG TACCGTAAGA	8100
AGCACCTGTA ATCATGATTG GTTTGTTCAA AAGTGGGTAAT ACCATAAG ACAACCAAGC	8160
AAGAGCGCTC ATCAAACAG CTGGAATAGA GTGATCATAC TCAGGAGTAC CGATAATAAC	8220
GCCATCTGCC TCTTCGATTT TAGCAGCAAT TTCCAATATT TCAGCAGGTA CTTGCTTGT	8280
AGCTGGTTG TTGAAGACAG GAATGGCCTT GATTTCAACA AGTTCAATT CAGCTTGTC	8340
AGTAAAGTGT TTTGCATGT ATTGAAGCAA TTGACGGTT GTAGAACGTT TTGAATTGTT	8400
TCCAACAATA GCAATAAGTT TAAACATGAG ATTTCTTTC TCTTTTACA TAATACAATT	8460
TTAAAATTCC ATTGAAACAG TTGCTCTAT AGAGTAGGAA TTCCTGAAGA ACAGCTTAGG	8520
TGGCCTCTT TATCGATGAG GATGACTTCG ATGCCCTCCA AACTTTCGAC TTGCCAGAGG	8580
ATAGAAGCAG GTCTTCTCC AAAGAGTCGA GTCGTCCAGA TTTGCCATC GACTGATT	8640
TCAGAGATGA TTGTTAGACT CGCTAGTTCC GTTCAACAG GATATCCTGT TTGACTGTCA	8700

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AAAATGTGAT GGTAATCTTG TCCATCGACG GTCAGGTGAC GTTCATAAAT GCCTGAAGTC	8760
ACGACAGATT TATTGACAAC AGGGATGGTC ATTAATGAT TTCCCCTAGG ATTGGCTGGG	8820
TCTTGAATCC CGATTGCCA TGGGTTATCC CCTCTTGCC GATTTTTTCC AATGGTCAGG	8880
ATATTCCTC CCAGATTGAT CAAGGCAGAA GTCACCCCCCT CTTTCCTAAG AAATTGGGCA	8940
ACCTTATCCG CACTGTATCC TTTGGCTAAA CAACCTAGAT CGATCTTCAT TCCTTTCTGT	9000
TTTAAAAACA CAGTAGAAGT AGAAGAATCT AACTCGATAC CATGAGGATT GATTAGAGGC	9060
AGCACCGATT CAATTCTTG AGGCTGGCG ACCTTGGCAT CTGAAAACC GATACGCCAG	9120
GTTTGAATTAA AGGGACCAAT GCTGATATTG AGGTGCTAG AGAGCGCTAG GCTATGCTCT	9180
AACCCAAGTG AAATCAGCTC AAACAGGTCT GGATGAACCG TGACGGGGC TATTCCCTGCT	9240
TGATAATTGA TTTCCATCAA CTCAGATTCT TGACTATTGG CGTTGAAGCG GTATTCAAGT	9300
TCTTTGAGCA AGTCAAAGGA TTTTTGGAGA AAGATATCGG CTTGCTCATC CACTAATGAA	9360
ATAGTGATAG TAGTCCCCAT TAGCCGTTCA GAATGTGAAC GAAGAGTCAG GCTACCAACT	9420
CCTTTCTCTT ATAGAAAATA AGTTGTAATA TCAAATAATC ATCTAAATTG AAGCCCTTAC	9480
ATTTCATTTT CATGTTATTA TAATACCATA AAGTTAGAAT TTTCACAAAC AAAATTTGGA	9540
AAAAGTCAG AAATATGCTC ATAAAATTCA TCAGGCTTGA AACAGGATA AATGGGGAAAT	9600
TATTTTGAT AAAAATGCT GAAATAATAG TACCCCCCTT GTAAACGCTA ACGGTAAATG	9660
GTATPACTAGT AAGGTAATT TAGAATGAAG GCAGGAAATT TTTATGAGTA AAATCGTTGT	9720
AGTCGGTGCT AACACCGCTG GTACAGCATG TATCAATACC ATGTTGGATA ATTTTGAAA	9780
TGAGAACGAA ATTGTTGTAT TTGACCAAAA CCTAACATC TCTTTCTAG GATGTGGAAT	9840
GGCTCTTGG ATTGGTGAAC AAATTGACGG TGCTGAAGGC TTGTTCTATT CTGATAAAGA	9900
AAAATTGGAA GCTAAAGGTG CTAAAGTTA CATGAACTCA CCTGTTCTTT CAATCGACTA	9960
TGATAACAAA GTAGTTACAG CGGAAGTTGA AGGAAAAGAG CACAAAGAAT CATAACGAAPA	10020
ATTGATTTTC GCTACAGGCT CTACACCAAT CTTGCCACCA ATCGAAGGTG TTGAAATTGT	10080
TAAAGGAAAC CGCGAATTAA AAGCAACTCT TGAAAACGTA CAATTGCTGA AATTGTACCA	10140
AAATGCTGAA GAAGTTATCA ATAAAATTTC TGACAAGAGC CAACACCTCG ACCGTATCGC	10200
CGTTGTTGGT GGTGGTTACA TCGGTGTTGA ACTTGCTGAA GCCTTTGAAC GTCTTGAAA	10260
AGAAGTGTC CTTGTTGATA TCGGTGATAC TGTCTTGAAAC GGTTACTATG ACAAAAGACTT	10320
CACACAAATG ATGGCGAAGA ACTTGGAAAGA TCACAAACATC CGCTTGGCTC TAGGTCAAAC	10380
TGTTAAAGCA ATCGAAGGTG ACGGTAAAGT TGAACGCTTG ATTACTGACA AAGAAAGCTT	10440

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TGACGTGGAT ATGGTTATCC TTGCAGTTGG TTTCCGTCGA AACACAGCCC TTGCAGGTGG	10500
TAAGATCGAA CTCTCCGCA ACGGTGCCTT CCTTGTAGAC AAGAAACAAG AAACATCTAT	10560
CCCAAGCGTT TACCGCTGTTG GTGACTGTGC GACTGTTAT GACAATGCTC GTAAAAGATAAC	10620
AAGCTATATC GCTCTGCTT CAAATGCTGT GCGCACTGGT AACGTTGGT	10669

(2) INFORMATION FOR SEQ ID NO: 58:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 7542 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 58:

CGCGCTAATA GATACTTTAT GATAGAATAA AGAACAAAGAT TGACAAGTAA GAGGAAACAT	60
TATGCAAAAT CAAACACTCA TGCAATACTT TGAATGGTAT CTGGCCCCACG ACGGTCAACA	120
CTGGACCGGT CTGGCTGAAA ATGCTCCACA CCTAGCTCAT CTGGGGATCA GTCACGTCTG	180
GATGCCACCA GCCTTCAGG CAACCAACGA AAAAGATGTC GGCTATGGGG TCTATGACTTT	240
ATTTGACTTA GGAGAGTTCA ACCAAAAAGG GACTGTCCGC ACCAAGTATG GTTTCAAAGA	300
AGACTATCTT CAAGCCATTC AAGCCCTTAA AGCACAGGGA ATTCAACCTA TGGCCGATGT	360
AGTTCTCAAC CACAAGGCTG CTGCCGATCA CAGGGAAAGCC TTTCAGGTTA TCGAAGTTGA	420
TCCCTGTAGAC CGTACAGTTG AACTTGGAGA ACCCTTCACC ATCAATGGCT GGACTAGTTT	480
TACCTTCGAT GGTGCCAAG ATACCTATAA TGGCTTCCAC TGGCATTGGT ACCACTTCAC	540
CGGTACAGAC TACGATGCCA AACCGAGTAA ATCTGGGATT TATCTGATCC AAGGGGACAA	600
CAAGGGCTGG GCCAACGAGG AATTGGTCGA TAACGAAAC GGAAACTACG ACTACCTCAT	660
GTATGCCGAC CTAGACTTTA AACATCCTGA AGTCATCCAA AACATCTATG ACTGGGCTGA	720
TTGGTTCATG GAAACGACTG GTGTAGCTGG TTTCCGTTG GATGCCGTTA AGCATATTGA	780
CTCTTTCTTT ATGCGCAACT TCATCCGCGA TATGAAGGAA AAATACGGTG ACGATTCTCA	840
TGTTTTGGT GAATTTGGA ACCCAGACAA GGAAGCCAAT CTGGACTATC TCGAAAAAAC	900
GGAAAGAACAC TTTGACCTTG TCGATGTTCG TCTCCACCAG AATCTCTTG AAGCCAGTC	960
AGCTGGCGCA AACTATGACC TTCGTGGCAT TTTCACAGAT AGCCTGGTTG AACTCAAGCC	1020
TGACAAGGCT GTGACTTTTG TCGACAACCA CGATACCCAA CGAGGACAAG CCCTTGAGTC	1080
TACCGTTGAA GAATGGTTCA AGCCAGCAGC CTATGCCCTC ATTTTGTAC GCCAAGACGG	1140
CCTTCCATGT GTCTTTACG GAGACTACTA TGGGATTCA GGGCACTATG CTCAAGAAGA	1200

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TTTCAAAGAA ATCCTTGACC GCCTCCTAGC CATCCGAAAA GATTTGGCCT ATGGAGAAC	1260
AAATGACTAC TTTGACCATG CTAACGTAT CGGTTGGTA CGTTCAGGTG CTGAAAATCA	1320
ATCCCCAATC GCAGTCCTTA TCTCAAATGA CCAAGAAAAC AGCAAGTCAA TGTTTGTCGG	1380
TCAAGAATGG ACTAATCAAAC CCTTTGTAGA TTTACTTGGT AACCACCAAG GTCAAGTTAC	1440
AATTGATGAG GAAGGTTATG GACAATTCCC TGTCAGCT AGATCCGTAA GTGTCCTGGC	1500
AGTCAATAAC ATCTAATAGC TCATAATAAC CAAGCTAGGT CCAAGCGGAT TTGGCTTTT	1560
TGTATTACACA AAAAGACCTA CCCAAATGGA TAGATCTTA CTTGATTACA ATTTACCTGC	1620
TACTGCATCC AACAAATTCTT GGATCTTAGG TTGGTTGCTT CCTCCTGCCA TGGCCATATC	1680
TGGTTTACCA CCACCAACGTC CATCGATGAT TGCGTCTAAAT TCTTTGACAA GGTTTCCTGC	1740
ATGAAGGTCT TTTGTCTTGC TTGCTACAAG GACATTGACT TTGTCACCGA TAGCGGCAAC	1800
TAGGACAAGA AGATCAGAGT AGTCTTTTG TTCCAGTTA TCTGCAAAAG TACGAAGGGC	1860
ACCGGCATCG GATACAGACA CTTGACTAGC AATGTAACGA TGACCGTTGA CTTCTTAAC	1920
ATCTTTGAAG ATATCGCCTG CGGTCGCAGC TGCGGCTTT TCTTTCAACT CAGCATTTTC	1980
TTTTTGAAAGT TGACGAAGTT GTTCTTGAAG TCCTTCTACC TTGTGAGGTAA CTTCTTGAC	2040
TTGAGGTGCT TTCAAGGTTG CTGCATAGC TTAAAGAGCA TCCTCTTGTT CACGATAGGC	2100
TTCAAAAGGCT TCCTTACCAAG TCACTGCCAA GATACGGCGA GTTCTGAAC CGATTCCTTC	2160
TTCTTGACA ATTTTGAAAGA GACCAATCTC AGAAAGTGTG TCAACATGAG TACCACCA	2220
AAGTTCAATA GAGTAGTCAC CGATAGTCAC GACACGAAC TCCCTGCCGT ATTTCTCAC	2280
AAAGAGGGCC ATAGCTCCCA TTCTTTAGC AGTGTCAATA TCCGTTCAA CTGCTTCAC	2340
TTCAAGTGCT TCCCCAATTT TCTCGTTAAC TTGCTGTTCA ATCGCACGAA GTTCTTCAGC	2400
AGTTACTGCT TGGAAGTGGG TAAAGTCAAA GCGAAGGAAT TCAACTTCGT TAAGAGATCC	2460
TGCCTGTGTT GCGTGGTTTC CAAGGATATT GTGAAGGGCA GCGTGAAGCA AATGAGTCGC	2520
AGTGTGGTTT TTCATGACAC GGTGACGGCG ATTGCTATCA ATTGCCAAGG TATATTCTTG	2580
GTTCAAGGCA AGCGGTGCAA GGACTTCAAC TGTATGAAGG GTTGTGACCAT TTGGGGCTTT	2640
CTGAACATTG GTCACAGTAG CCACAACTT ACCTGACTCA TCCAAGATTT GTCCGTAGTC	2700
AGCTACCTGT CCACCCATT CAGCATAAAA TGACGTTCC GCAAAGATAA GAGAGGCAGT	2760
TCCCTCTGAA ACAGCTCCTA CTTCTGCATT GTCAGCAACG ATAGCTACCA ATTTAGAAGA	2820
CAATTGGCTA GCATTGTAGT TGAAGACACT TTCTACAGTG ATGTTTGAA GAGTTTCATT	2880
TTGCATACCC ATTGAGGCCAC CCTTGACAGC TGACGCACCC GCGCGTTCTT GCTGTTCTT	2940

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CATGGCTGCT	TCAAAACCTT	CACGGTCTAC	AGTCATACCA	GCTTCTTCAG	CGATTTCTTC	3000
AGTCAATTCA	ACTGGGAACC	CATAAGTATC	ATAGAGTTG	AAGACATCTG	AACCAGCGAT	3060
AACAGATTGA	CCTTTTCTT	TCAAGTCTGC	TACAATGCC	TGGGCAAAGT	GTTGACCTGA	3120
GTGAAGGGTA	CGGGCAAATG	ATTCTTCTTC	GCTCTTAACG	ATTTTCTCAA	TAAAGTCACG	3180
TTTCTCAAGC	ACTTCTGGGT	AGTAGCTTTC	CATGATTTT	CCAACAGTTG	GAACCAATT	3240
GTAAAGAAA	GGCTCGTTGA	TACCCAATT	TTGACCATGC	ATAGAAGCAC	GACGGAGAAG	3300
ACGACGAAGA	ACATAACCAC	GACCTTCATT	TCCTGGAAGG	GCACCATCAC	CGATAGCAA	3360
TGAAAGAGAA	CGAATGTGGT	CTGCGATAAC	CTTGAAGCTC	ATGTTGTCGC	CATCTGGTC	3420
ATAAACCTTA	CCAGACAATT	TCTCGACTTC	ACGGATAATC	GGCATGAAGA	GGTCCGTTTC	3480
AAAGTTGGTC	TTAGCCCCCT	GGATAACGGC	CACCAAACGC	TCCAAACAG	CGCCCGTATC	3540
AATGTTCTTA	TGTGCAATT	CCTTGTATTTC	GCTACGAGGA	ACAGCAGGGT	CTGCGTTAAA	3600
TTGTGACAAA	ACGATGTTCC	AGATTCAAT	ATAACGGTCG	TTTCAATAT	CTTCTGCAAG	3660
CAGGCGAAGA	CCGATATTTC	CTGGGTCAA	GGCTTCCCCA	CGGTCAAAGA	AGATTCTGT	3720
ATCTGGTCCA	GAAGGTCGG	CACCGATTTC	CCAGAAGTTG	TCCTCAATTG	GAATCAAGTG	3780
ACTTGGATCC	ACTCCCACCT	CAATCCAGCG	CTTGTAAAGA	TCTTTATCGT	CTGGATAGTA	3840
GGTCATGTAA	AGTTTTTCAG	CAGGGAAATC	AAACCATTC	GGGCTTGTCA	AAAGCTCATA	3900
AGCCCAAGTG	ATAGCTTCGT	CACCGAAGTA	ATCCCCGATA	GAGAAGTTCC	CCAGCATTTC	3960
AAACATGGTA	TGGTGACGCG	CGGTCTTCCC	TACGTTTCG	ATGTCGTTGG	TACGGATAGC	4020
CTTTGGGCA	TTGGTAAATAC	GTGGATTTTC	AGGGATAATG	GTCCCGTCAA	AGTATTCTT	4080
AAGGGTTGCT	ACCCAGAGT	TGATCCACAA	AAAGAGTTGGG	TCATTTACAG	GAACCAAAC	4140
TACTGATGGT	TCTACTGAGT	GACCTTTGGT	CGCCCGAGAA	TCAAGCCACA	TTTGGCGTAC	4200
TTGTGCACTA	GATAGTTGTT	TCATATTGTC	TCTTATTCA	CTTGTGTTAAT	CTGATTGGCT	4260
TTCCACCATG	TCCACATAGT	CAATCGCGAC	ACAGAGGGAA	ATGACTAGGT	CTGCATAAGC	4320
GTCTTCAAGA	ACCGTTACGG	TATAGGTAGA	AGTCAGATGG	AAGAGTTCC	TCTTAATTTC	4380
CGCAATCAAC	TGATCGCGAT	CATCCAGAA	TTTGAAATT	AAATCCCAGA	TATTGCCCTC	4440
GATACGAAGA	CCTAGATTAT	CAAACTCATA	CTTATCTCGC	CAGAAGGTCA	ACTTCTTACG	4500
AATGACAAAA	CTCGAGCCAT	CCCGAAGCTG	AATTCAAAA	CGAGGAAGCA	AGGTCAAGAT	4560
TTCTTTACTA	ATCTCACTGA	CTTGTTCACC	AGCCGCATCA	TAGATGGTAA	AGGTTTTAGG	4620
AATCTAAAAA	AATGATCCCT	CCACCTGATA	GGCAATTCT	CCCCTGTCA	CCTTGATAGC	4680
GAAGCGTTCG	CCTCCAAGAC	GAAACTTTG	TTTGACAAGA	AATGTTTCA	TCAACACCTC	4740

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CAAAAATCAA AAGACAAGCT CATATCACGA AGGGCGAAAA ACCGCGGTAC CACCTTCATT	4800
CAATGAAC TT GTCATTCTCT TGTCTTATG CAATTGTATG ATTGAGTAGC ATGACTTCCT	4860
AGCTTAGATG GCTCCGAGCA CGGCCATTTC TCTGGACTAA GACAAGTGAA AATCAATTCT	4920
CAACTTCTT ATTATAACGT TTTTTTAAGC TTGCGTCAAC TGGAAATGAT CTCCGTTGAA	4980
TTAGACCAAT TCCCTACATC TCTGATTACT TTTTCAGGAT ATATTTTTTC TTACTGCCT	5040
TTTTCTTTT ATCCCAAATT TTCATATTAC TAAACACAGC TACTAGAATA TTTCCAAATA	5100
TAAAGGTGCC TATCACCCAA TATATGGACT CAGTTGTTAG GTATTGTCGA TCCAAGCCAT	5160
CCTTTAAATG GAATAGTATA GCAGTTGGT TAACAATCAT AAAGGTTGGC CAGAAACTTT	5220
TTTTGAAAAA AGTAGACATT TTCATTATTT GTTGCGCTT TCTGTAAGGT TAATACTCAA	5280
AAAAAAATCAA AAAGCAAAC AGGAAGCTAG CCTCAAGCTG TACTTGAGTA CGGCAAGGCA	5340
ACGCTGACGT GGTTGAAGA GTATAGGCTT AGTATACTAC TAGGCAAGCA AATAAACAAA	5400
TAAACAACTA GAATAGAAAA AGATAGGGCT CTAAAAACTG ACTTCTATTTC CTTAAAACG	5460
AACCAGCTT ACTGATTCTG CTTCTTACGT TTATCTCCTA CTTCCGATAC ATTTTAAACT	5520
GTAGGAAGAG GTCGCTATAT TTCCCTGTCC ATTTATGGTC AAATTTCTCA TAAACTTCTA	5580
GGTCTTCAT GGTTCAACA TCGGGATAGA AGGCCTTATC TTCCCTTGTT TCCTCTGGGA	5640
GCAATTCCTT CGCTCGTAGG TTTGGTGTG AATAGCCGAC ATACTCCGCA TTTTGGAGAG	5700
CATTTTCAGG TTTCAACATA AAGMTGATAA AGGCATAGGC TGAGTTTTGG TTTTTAACTG	5760
TTTTGGGAAT GACCATAATTG TCAAACAAA GATTGCTGGC CTCTGTCGGT ACCACATAAC	5820
GTAGATTTTC ATTTTTCT AACATTTGGC TGGCTTCACC AGAGAAGGTC ACGCCGATTG	5880
CAACATTATT CTGAATCATA TAGCCCTTCA TCTCGTCCGC AACGATAGCC TTGATATTG	5940
GAGTCAGTTT GTAGAGCTTA TCCACTGTCT CTTCCAAGTG CTGCAGATCC TTGGAGTTGA	6000
GGCTGTAGCC GAGGAATTG AGTCCTAGTC CCAGCACCTC ACGCGCCCCA TCAAAGAGCA	6060
TGATAGAATT CTTATACTCC GGCTTCCAAA GGTCATCCC ATGCTCAGGC GCTTCATCTA	6120
CCATGGTTTC GTTGTAGACA ATTCTTAAGG TTCCCCAGAA GTAAGGGATG GAGAATTAT	6180
TACCTGGTC AAAGGACTGG TTGAGAAACT CTGGTCCGAT ATTTTCGATT CCTTCATTT	6240
TTGAATAATC AAGCGGAACC AAGAGGTCTT CGTCCTTCAT CTTGTTAAC ATGTATTAC	6300
TTGGAATGGC AATATCGTAG GTCGTTCCAC CCTGCTTAT CTTAGTGTAC ATGGCTTCGT	6360
TGGAGTCAAAGTCTCGTAC TGAACCTGAA TTCCCTGTTTC TTCTGTAAAC TGAGTCAAGA	6420
GTTCAAGGATC GATATAGTCT CCCCAGTTAT AGATAACCAA TTTTTGACTA TCTCGACTAT	6480

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TGATTTTACT ATCTAAATGA GTCGCAATTG	6540
CCACAAAGAC AAGGATAATC GCTGCAATTG	
CTGCTAAAAA TGAATAGATT TTTTCATGC TTGCTCCTCC TTCTCACGAG AGATAAAAGTA	6600
ATAAACCTACA ACTAGGATAA TACTAAAGAG AAAGACTAGA GCAGACAGGG CATTGATTTC	6660
TAAGGAAATC CCCTTGCAG CACGAGAGTA AATCTCGACT GATAGGGTTG AAAAGCCATT	6720
TCCTGTTACA AAGAAGGTCA CGGCAAAGTC ATCTAACGAA TAGGTGAAGG CCATGAAATA	6780
ACCAGTAATG ATAGACGGAG TCAGGTAAGG AAGCATGATT TCCTTGAACA TCTGAAATTG	6840
ACTAGCTCCC AAGTCATAGG CCGCATGAAT CATGTCGCCA TTCATTTCT TGAGTCGAGG	6900
CAAGACCATC AAGACCACGA TAGGAATGGA GAAGGCCACG TGACTAGATA GAACGGTCAA	6960
AAAGCCAAGT GAAAACTTGA GTTGGGTAAA GAGAATCAAG AAGCTAGCAC CAATCATAAC	7020
GTCAGCGCA ACCATGAGGA TATTATTGAG TGATAGAAAG GCTCTTGGT ATTTCTTACG	7080
AGACTGGTAG ATGTAATGG CACCAAAAGT CCCGATAATG GTCGCTATCA AGGCTGATAG	7140
GAAGGCCAAG AAAATGTCT GAGCCAAAT CAGCATGAGT CTCCCATCTC CAAACATGGT	7200
TTCAAAGTGA GTCCAGCTAA AACCTGTAAA GCTATTCTATA TCATCACCAAG CATTAAAGGC	7260
ATAGCCAATC AAGTAAAAGA TAGGCAGGTA GAGGACCAAGA AAGACCAGTC CCAGATAAAG	7320
GTTGGCAAT TTTTCATCG TTCTCTCCTT TCCTTAGTCA CCCACATGGT GATGAACATG	7380
GTCAGGATGA GAATCACACC GATGGTTGAA CCCATACCAT AGTTGTCATT GGTTAGAAAA	7440
TTCTGCTCAA TAGCCGTCCC CAAGGTGATA ACGCGTCCC ACCAATCAA CGGGTCAGCA	7500
TGAAGAGACT CAAACTTGAG ATAAGACCG ACTGAACCCC GG	7542

(2) INFORMATION FOR SEQ ID NO: 59:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 9223 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 59:

AAAACCAAAT TCCGGTATTT TAACCTATGC TGAAATACC ATGAAGTCTG TCATGACAGA	60
TCAGGTCTAT AACATTAAGG TTGAGACAGA AAATGGAAAT TATGTTGGTG AAGCTAGCCA	120
TGTTTGTC CTTTGACAA ATTACTTCGC TGATAAGAAA ATCTTTGAAG AAAACAAGGA	180
CGGCTATGCC AACATTTGTA TTCTGAAAGA TGCCCTCTATA TTCTCCAAT TATCCGTCAT	240
TCCTGATTTA TTAAAAGGGG ATGTTGTCGC AAATGATAAT ATCGAGTATA TCAAAGCGCG	300
TAATATTTAA ATCTCTTCAG ATAGTGAATT GGAGTCAGAT GTTGACGGAG ATAAATCAGA	360

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TAACCTACCT GTAGAAATCA AAGTCCTAGC TCAGCGAGTA GAAGTATTTT CAAAACGAA	420
AGAGGATTAG TATATAGAGA AAGCCTTTT TAAGGCTTT TGTATACTTT AAAAGATAGT	480
TCCTTTAACCA ACGGACATTC CTTGCAAATA GTTTACAAA AATAGTATAC TGGATTCAATT	540
GAGTTTGAAA ACGTTTGCCTT AAAATTGAA TGAATACTTT AGGAGACAAA TTGATGGAAT	600
TGAGTGCTAT TTACCATAGG CCTGAGTCGG AGTATGACTA TCTTTATAAG GATAAGAAC	660
TCCATATTGCG AATTGAACT AAGAAAGGGG ACATTGAAAG CATCAACTTG CACTATGGGG	720
ACCCCTTTAT CTTTATGGAG GAGTTTATC AGGATACAAA AGAAATGGTC AAGATAACTT	780
CTGGTACCTT ATTTGACCAT TGGCAGGTTG AAGTGTCACT TGACTTTGCA CGTATCCAGT	840
ATCTCTTGAA GCTCAGAGAT ACAGAAGGTC AAAATTTT GTATGGCGAT AAAGGGTGTG	900
TGGAAAATTC TCTAGAAAAT CTTCATGCAA TTGGGAATGG ATTTAAGTTG CCTTAGCTTC	960
ATGAGATTGA TGCCTGCAAG gTTCTGACT GGGTTCAAA TACGGTATGG TATCAGATAT	1020
TTCCTGAAAG ATTTGCCAT GGCAATGCTC TATTAAACCC AGAAGGGACT TTAGACTGGG	1080
ATTCATCTGT CACACCTAAG AGCGATGATT TCTTTGGTGG TGATTTACAG GGGATTATTG	1140
ATCATATGAA TTACTTGCAA GACTTGGTA TTACTGGACT ATATCTTGT CCCATCTTG	1200
AATCTACAAG CAATCACAAG TACAATACGA CAGATTACTT TGAAATGAC CGTCATTTG	1260
GAGACAAGGA GACCTTCGG GAACTGGTGG ATCAAGCGCA TCATCGTGGC ATGAAAGTCA	1320
TGCTGGATGC GGTATTTAAT CATATTGGTT CGCAATCTCT TCAATGGAAA AATGTCGTCA	1380
AAAATGGTGA ACAGTCTGCT TATAAGGATT GTTCCATAT TCAACAATTC CCAGTGACAA	1440
CTGAAAAGCT AGTTAATAAG AGAGACTTAC CCTATCATGT TTTGGTTTC GAGGACTATA	1500
TGCCTAAGCT AAATCACGCC AATCCAGAGG TCAAGAATTA TCTTTAAAG GTTGCAGCT	1560
ATTGGATTGA AGAGTTTAAT ATCGATGCTT GGCGTTGGA TGTGGCTAAT GAGATTGACC	1620
ATCAGTTCTG GAAGGATTTT CGTAAGGCAG TTTTAGCTAA AAATCCGTAT CTTTATATCC	1680
TAGGAGAAGT CTGGCATACA TCTCAGCCTT GGCTAAATGG AGATGAGTTC CATGCCGTCA	1740
TGAATTATCC TTTATCTGAT AGTATCAAGG ACTATTCTT ACGAGGAATT AAGAAGACAG	1800
ACCAGTTCAT CGATGAAATC AATGGAGAGT CTATGTATTA CAAGCAGCAG ATTTCAGAGG	1860
TCATGTTAA TCTCTTGGAT TCACATGATA CAGAGCGAAT CCTGTGGACG GCCAATGAAG	1920
ATGTTCAACT GTTAAATCA GCCTTAGCCT TTCTCTTTT ACAAAAAGGA ACACCGTGCA	1980
TTTATTACGG AACCGAGCTA GCCTTGACTG GAGGACCAGA TCCAGATTGT CGTCGTTGTA	2040
TGCCTGGGA ACGTGTATCA AGTGACAATG ATATGCTGAA CTTTATGAAG AGGCTGATTA	2100

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AAATTCCGAA	ATACCGGTCA	GTAATCATT	CGCATGGCAA	GTATAGCCTT	CAAGAAATCA	2160
ACTCTGATCT	AGTAGCTCTG	GAATGGAAAT	ACGAAGGACG	GATCCTAAA	GCAATATTCA	2220
ACCAATCAC	AGAAGATTAT	CTTTAGAGA	AAGAACGAGT	AGCACTAGCA	AGCAATTGCC	2280
AAGAATTGGA	TAATCAGCTT	GTCATCTCTC	CAGATGGATT	TATGATTTTC	AAAAAACTAG	2340
TTGATGAAGA	TTATGGTACA	TTTCATACCT	TATATAGTAT	AATAAGGCTA	GTACTAAAC	2400
TTGTAAGGA	GAACCTAAAT	GAATTGTTAGA	GGACATGAAA	CAAGACAAAG	AATTGTTAGA	2460
GATTTTGAAG	TTCAGCCTAA	AGCACATATT	AAGCTGTTAG	CAAATCAACA	AAAACATAGT	2520
GATGCAGGAG	CAACTATTGA	AGATGAATAT	TATGTATT	TCGCTGAGAG	AAAAATTGAT	2580
GGCAAGAAGG	AAAGTTATTCA	GTGTTGCATG	GGTGCAGCAA	GGGATTTTTT	AGAACTAATT	2640
AATCACAAAG	GGCTACCTCT	TTTAAATCCG	CTTGTAGGTG	ATTCTCATGT	AAATAATAGA	2700
CAAGAATATG	ACAATACAGG	GAGTGGAAAT	TTATAACCTG	AAAAGTGGAA	TGAAACTGCC	2760
AAGCAGCTT	ATAATGCTAT	AATGTTGGTTG	ATTATTTTAT	GGAATGCTAA	GCCGGATACA	2820
CCTTTATTTA	ATTTAAAGA	CGAAGTAATT	AAGTATAAAA	CATATGAGCC	TTTGAAAGC	2880
AGTATAAAA	GAGTAAATAC	TACTATAAG	AATGGTAGTA	AAGGGAAAAC	TCTGACTGAG	2940
ATGATTAATG	GCTACAGAGC	GGATAACGAT	ATTAGAGATG	AAATTTGTAA	CTTTAATATT	3000
CTGAAAAATA	AAATTCTGTGA	TATGAAAAAC	CAACAAGGAA	ATACAATGGA	ATCTTACTTT	3060
TAGTTATTGT	TGAATTGTTGG	GTATTCTATA	AAATATCCTA	ATTGAGATTT	AAATAGTAGA	3120
CTATACAAATA	TAGTTAAAT	ATCAGTAAAAA	ACAACACTTT	ATTGAGGTAT	TGGATACGCT	3180
TTGCTAATAG	CCTAATAATC	ACATGTGGAG	TGTTGCTACA	ACGAAAAAGG	TGATAATCCT	3240
TGATTTCAAG	CTATTTTATA	AGCATTGTTGT	CTTGTAGAT	AAAGGCAATT	TTGACAATAA	3300
AAATCCTAAA	AGGTGAATCG	TTATAGATGT	ATTGTTAGAT	ATCGTTGCG	CATCGAAAAA	3360
ATTAATACAA	GAATAAAAT	TTATAGCTCT	TTAGGTACT	TTTATAGAAG	AAAGTTTAG	3420
GATAGAAAAA	CAAGAAATAA	CGCACCATTT	TTGGTGCCTT	ATGCTTTTTT	ATGCTATAAT	3480
GGATTTATAA	AAATAAAGGA	GTTTGCTATG	ATTGGAAAGA	ACATAAAATC	CTTGCCTAAA	3540
ACACATGACT	TAACACAACT	CGAATTGCA	CGGATTGAG	GTATTCACG	AAATAGTCTG	3600
AGTCGTTATG	AAAATGGAAC	GAGTTTCAGTC	TCTACCGAAT	TAATAGACAT	CATTGTCAG	3660
AAGTTTAATG	TATCTTATGT	CGATATTGTA	GGAGAAAGATA	AAATGCTAA	TCCTGTTGAA	3720
GATTATGAAT	TGACTTTAAA	AATTGAAATT	GTGAAAGAAA	GAGGTGCTAA	TCTATTATCT	3780
CGACTCTATC	GTTATCAAGA	TAGTCAGGGA	ATTAGCATTG	ATGATGAGTC	TAATCCTTGG	3840
ATTTTAATGA	GTGATGATCT	ATCTGATTG	ATTCATACGA	ATATCTATCT	AGTAGAAACT	3900

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TTTGATGAAA TAGAGAGATA TAGTGGCTAT TTGGATGGAA TTGAAACGTAT GTTAGAGATA	3960
TCTGAAAAAC GGATGGTGGC CTAATGGAAA TCCAAGATT AACTGATAGT GAATTCAAAC	4020
ATGCTTCTGC AAGGAATCTT CGTTCACTGA CAAGAGGAAA AAAGTCCAGT AAGCACACCA	4080
TAGCGATTTT GCTTGGAGGG CAAAGTGGTG CCGGTAAGAC TACAATTCA CGTATTAAC	4140
AGAAAGAATT TCAAGGAAAT ATTGTTATCA TAGATGGTGA TAGTTTCGT TCTCAGCATC	4200
CACACTATTT AGAACTGCAG CAAGAATATG GCAAAGACAG TGTAGAATAT ACCAAAGATT	4260
TTGCAGGAAA AATGGTAGAG TCTTTAGTAA CAAAATTGAG TAGTTGAGA TACAATCTT	4320
TGATAGAGGG AACTTTACGA ACAGTTGATG TTCCAAAGAA AACAGCACAA CTCTTGAAA	4380
ATAAGGGATA TGAAGTACAA TTGGCCTTAA TTGCGACAAA GCCTGAATTG TCGTATCTAA	4440
GTAACCTTAT CCGTTATGAA GAACGTGTACA TTATCAATCC AAATCAAGCA CGCGCAACTC	4500
CAAAAGAACCA TCATGATTC ATTGTAATTC ATCTAGTTGA TAACACACGA AAATTGGAAG	4560
AACTAGCTAT CTTTGAAAGA ATTCAAATTT ACCAACGAGA TAGAAGTTGT GTATATGATT	4620
CAAAAGAAAA TACAACCTTC GCAGCAGATG TTCTTCAAGA GTTACTCTTT GGGGAGTGG	4680
GTCAGGTAGA GAAGGGAGATG TTGCAGGTGG GGGAAAAGAG ACTTAATGAA TTACTTGAAA	4740
AATAAACAAAT TGATATTTTT AGGAGAAATAG AAATGAGAGG GTTTAATAAC AAGATAAAAGT	4800
CTGTTATCA AGAACTAACAA ATTCCAAAG AGAAATTCCG TAGCTTTCAC AAGACTTTAA	4860
TTCATTTCCA TACACCTGTT TCTTATGATT ACAAGCTATT TTCTAATTGG ACTGCAACGA	4920
AATATAGAAA AATTACTGAA GATGAACATAT ATGATATATT TTTTGAAAAT AAGAAAATAA	4980
AAGTTGATAA GACAATTTT TTTAGTAATT TTGATAAGGT TGTTTTCT AGTTCAAAG	5040
AATATATTAG TTTCTTATG TTAGCAGAGG CAATCATAAA AAATGGAATA GAAATAGTTG	5100
TAGTAACGTGA TCATAAACTACT ACCAAAGGTA TTAAGGTT ACAATGGCA GTCTCAATCA	5160
TAATGAAAAA TTATCCGATT TATGATATAC ATCCTCATAT TTTACATGGA GTAGAAATTA	5220
GTGCAGCAGA TAAATTGCAAT ATTGTATGTA TATATGATTA TGAACAAGAA TCATGGGTTA	5280
ATCAATGGTT AAGTGAATAT ATTATAAGTG AGAAAGATGG AAGTTATCAA CATTCACTGA	5340
CTATAATGAA GGATTTCAAT AATCAAAAAA TAGTTAACTA TATTGCTCAT TTCAATAGTT	5400
ATGACATTTT GAAAAAAGGT TCTCACTTAT CAGGTGCATA TAAACGAAAA ATTTTTCTA	5460
AAGAAAATAC ACGATTTGG AGTTTAATAT TAACTCGAAA GAATCTTCGC AACAACTTGA	5520
TATTCTCTAT AAAGAAGTTG GTGTATTAAG TTTGGGACAA AAAGTTGTAG CCATGCTTGA	5580
TTTTTATTA GCATATAGTG ATTATTCTAA AGACTTCAGA CCATTGATTA TTGATCAGCC	5640

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TGAAGACAAT	CTAGACAATC	GTTATATTAA	CAGGCATTAA	GTTCAGCAGT	TTAGAGATGT	5700
GAAAGCTCAA	CGTCAAATTA	TTTAGCAAC	ACATAATGCT	ACAATTGTA	CAAATTCTAT	5760
GACAGATCAA	GTTGTTATTAA	TGGAGTCAGA	TGGAGTTAAC	GGATGGATTG	AATCACAGGG	5820
ATATGTTAGT	AAAAAATATA	TAAAAAATCA	TATCATCAAT	CAATTAGAGG	GAGGAAAAGA	5880
TTCCTTCAAG	CATAAAATGT	CTATATATGA	GACGGCTTTA	TCAGAGTAGA	GTCAGAAAAA	5940
GTAGGGTACA	AATTAGCCT	ACTTTTTCT	TTGTCGACAA	GGCATAGTGT	ACATCTGAGG	6000
TCCAAGTCCT	CTGTGGATAT	TTGCTGCAGA	TGAAACCAAT	AGCGACTCCT	AAGCCTGAAT	6060
ATCGTGAGGT	AGGGGGATA	GGAAGGAATT	AGCGAAATCA	AGGTTCTACA	AACAGAATCG	6120
TGACTTGAAAG	CCATATATAG	CGGATGAGGA	ACTCTAAAT	CCAAATAGGT	GTCGTAACCT	6180
ATATACGTAA	ATTACGAGAG	AAAACAGGA	AAGATGTACG	GCTTATTCCG	TGAGCGTTA	6240
GGACGTAGTA	CAACGAATCA	TGGGAGTCAG	CTGAACACAT	AGTATTGAA	AAATTTCTGT	6300
AATGGAATAG	GAGCGAAGAA	GTGAAACAATT	AAATGAATAC	CTCTCTAATT	AAATTTGTCA	6360
ATTCTAAATTC	CTGGTATGAA	AAGACAGTGA	CCTGAAAATG	TAAACGATGG	GAGCTGATCA	6420
TAAATATAGG	ACGGTACATG	CACTGGTGT	AGAGATTAGT	CCTTACTTGA	TTTGTGATAA	6480
CTTCCCCAAA	TTTCTTCTGC	TATACTTTTC	TCAACTTTA	AAAATCCAAC	TAAGAATT	6540
ACCTGGGGGT	TTGGGGCGGG	AGCACTAAGT	TATCTTATCG	TTAGCTGTCA	AAACTGGTAG	6600
GTTTGATAG	GCTGGCGATA	TGATTTTG	GATATTGTGG	ACACAATATC	TGAGCTCGCA	6660
AAGCCTTACA	AGAATGAAAA	TCAGTTGTTG	AAAAAGTGTA	CTGACATTGT	ATGGTAGCTC	6720
ACATTGTCAG	TACAAGTATT	TTGGAAAGGA	AGTAGCAGTA	TGAAACGAGA	TGTGGTGTAT	6780
ATTCGAAAC	AATTCTGTTT	AACAGAAGCA	GAAGAAAAGC	AAATTCTAGC	TTTGATGAGA	6840
GAGCGGGGAG	AGACTAATT	CTCTGATTTT	CTTCGTAAA	TTTACTTTC	CTCTGATT	6900
CAAAACAGA	TGGAGACATG	GTTCGCCCTC	TGGCAATCCC	AAAAACTAGA	ACAAATCAGT	6960
CGTGACGTTC	ATGAAGTTT	AATCTGGCA	CAGTCAGAAC	GTCAAGTCAC	CCAAGAGCAT	7020
GTATCTATT	TCTAACGTG	CGTGCAGGAA	TTGATTCAAG	AGGTTGCAAA	CACCACCC	7080
CTCAGTAAAG	AATTCTGTGA	GAAGTACATG	AGGTAAGCAC	ATGGAACATC	GTTACCGAAC	7140
CAATCTCAAG	AAAGTGT	TGTCTGATAG	TGAGTTGAAC	CAACTAAATA	AAATATCGA	7200
TCAAAGTGGT	TGTAATCCT	TTTCTGAATA	TGGAGACGA	ACTCTACTCG	ATCTGGTAT	7260
GAATTTATC	ACGATTGACA	CAAACGGTTA	CCAAGTTA	GTGTTGAGT	AAAGAGGAT	7320
TGGCAATAAT	ATCAACCAGA	TTGCTCGAAG	TGTTAATCAA	TCTCAGTTAA	TTTCTGGTGA	7380
AGAATTGCA	GAGTTGAAAA	AAGGAATTGG	TGAATTGATA	AAAGAAGTTG	ATAAGGAATT	7440

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TAATCTGCAA GCGCAGAAGC TAAAGGAGTT CCATGGTCAT CACTAAACAC TTTGCCATT	7500
ACGGAAAGAG TTACCGCAGA AAGCTTATCA AGTACATTCT CAATCCTGAG AAAACCAATA	7560
ATCTTGCCCTT GGTCGCGAC TATGGCATGA AGAATTTCT GGACTTTCT AGCTATGAGG	7620
AAATGGTGCA GATGTATCAT GAAAATTTCA TCAGCAACGA TACGCTTAC GATTTTCGCC	7680
ACGACAGGAT GGAAGAAAAT CAACGAAAAA TACACGCTCA CCACATCATT CAGTCTTCT	7740
CGCCAGAGGA TCATATCACT CCTGAACAAA TCAATCGGAT AGGTTATGAG ACTGTGAAGG	7800
AATTAACTGG TGGCAAATTT CGTTTATCG TTGCGACCCA TGTTGATAAA GACCACCTGC	7860
ACAATCACAT CATTATCAAT TCACTAGATA GCAATTCTGA CAAAAAGCTC AAGTGGACT	7920
ACAAGGTGGA GCGAAATCTT CGCATGATTT CTGACCGTTT TTCTAAAATC GCAGGGCTA	7980
AAATCATTGA GAACCGCTAT TCTCACCAGC GGTATGAAGT CTATCGTAAG ACTAATCACA	8040
AGTATGAACT CAACGAGCGA CTCTATTTTT TGATGGAACA TTCTAGGGAC TTTGAGGATT	8100
TCAAAAAGAA TGCTCCGCTA CTACATGTGG AGATGGATTT CCGTCACAAG CATGCCACCT	8160
TTTTTATTAC GGACTCAACT ATGAAACAGG TGGTGCCTGG CAAGCAACTC AATCGCAAGC	8220
AGCCTTACAC AGAAGAATTT TTTAAGAACT ACTTTGCCAA AAGAGAAATA GAAAGTCTCA	8280
TGGAATTTTT ATTGCTGAAA GTTGAGAATA TGGATGATTT ACTTCAGAAA GCAAAACTTT	8340
TTGGACTAAC TATCAATCCT AAACAAAAGC ATGTTCTTT TCAATTGCA GGAGTGGAGG	8400
TAAAGGAGAC AGAGCTAGAC CAGAAAAATC TTTATGATGT AGAGTTTTTC CAAGATTATT	8460
TTAAAAATAG AAAAGATTGG CAAGCTCCAG AACTGAGGA TTTCGTTCAA CTTTATCAAG	8520
AAGAAAAGTT ATCCAAAGAA AAAGAACTTC CAAGCGATGA GAAGTTCTGG GAGTCCTATC	8580
AAGAGTTCAA GAGTAACAGA GATGCCGTTA ATGAATTGAA GGTGGAGTTG TCACTCAATC	8640
AAATTGAAAA AGTAGTGGAT GATGGAATTT ACGTCAAGGT CAAGTTGGT ATTCTGTCAGG	8700
AGGGACTTAT CTTTGTGCCG AACATGCAGC TTGATATGGA AGAGGATAAG GTGAAGGTTT	8760
TCATCAGGGA ACCAGCTCC TACTATGTCT ACCACAAAGA CGCTGCCAG AAAAATTGTT	8820
ATATGAAAGG TCGAACCTTA ATTAGACAGT TCAGCTATGA AAATCAAACC ATTCCATTAC	8880
GCAGAAAAGC GACAGTCGAT ATGATTAAAG AGAAGATTGC GGAAGTGGAT GCTTTGATTG	8940
AACTGGAAGT AGAAAATCAA TCTTATGTCA CGATTAAGA TGAGTTAGTG CATGAACCTAG	9000
CAGCGTCTGA ATTGAGAATC AATGAGTTGC AAGAACGAAT GTCAACCTTG AATCAAGTAG	9060
CAGAATATCT ACTGGCTTCA GTTGAAAGTA AGCAAGAAAT GAAATTAAT CTTTCAAAAC	9120
TGAATATAAC TGAGAAATAC AGTGCTAATA TTGTTGAGAA AAAATTGAAG AGCCTGGGA	9180

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ATCAACTGGA ATTGGAAAGG GGCAGGTATG AAAAGATGGT AGT 9223

(2) INFORMATION FOR SEQ ID NO: 60:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 6827 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 60:

TCTGCTGGCT ACCATCATCT GACTGGGCA AGACCAAAGT CTTAGTTACA ACTGTATTCT	60
TCTCAGCATT TTCAATAACT GGCAATGCCG ACTGAAGCGT ATCTTTTCT GTTTTTGTAG	120
CTGGTCCAGT TTCTTTTTC TGTCGCAAC CAACCAGGAC AAAAAGGAAA GCTAGACTAA	180
CAAGAACTAT TTTTTTCATT TCTTTCTTCT TTCTTTTGA AATTAAAAATA GAATAAGACT	240
GGGAAGTGCT CCCAGCCTTG ATGTTTATAG AGCTGCACGC AAACGTGCTT CTGCATTTTC	300
TACATTACGG ACAGAGCGTG GTAGGAAGGC ACGAATATCG TCTTCCTTGT AGCCAACTTG	360
CAGGCCTTT TCATCTACAA GGATTGGGCT CTTTAAAATT CTCGGTGTTC CCATAATCAG	420
ATTGAGAACT TCATTGACAC TCAAATCTTC AATATCCACT CCAAGGGCTT TGGCATAGCG	480
ATTTTAGAC GAAACGATGC TGGCTATTCC GTTATCTGTT TTGGTTAGAA TATCCAGTAA	540
TTCTTCTCTC GTAATTCCCTT CTTTACCAAG GTTTGTTCT TTATAACTTA ACTGGTGGGC	600
ATTGAGCCAG GTTTTGCTT TTTACAGCT AGTACAACCTT GAGACTGTAT AAATTTTAAT	660
CATGTACCTA CCCCTTTCGC TACATGTTAC TATCAGTTA GTCTATTATA CCATAAAAAA	720
CATCCGACTT GCGACCTATT TTTAATTTTT TTGACTTTT TCGTCATTT TCGTACTTTT	780
TTCTTGACAA ACAACTAAAT GACTATCAAC TCTTTGGAG CTAGGGTCAA TAATTACAA	840
CCTGTCCTCG TAATCAGGAT ATCATCCTCG ATACGAACGC CATATTGCC TTCGATATAG	900
ATACCTGGTT CATCGTCAA GGCCATACCT GTCTTAATAG TTTCTGTAGA AGTCTGACTA	960
AAGTAGGGTT CCTCATGGAT ATCCAGACCA ATACCGTGGC CAATGCCGT AGTAAAGTAG	1020
TCACCATAAC CTGCCTCAAT GATAATATCA CGAGGGATTT TGTCAAAGTC ACGGAAACCT	1080
AAGCCTGCCT TAGCTTGGTC AATCAAGGCT TGTTAGCTT TTAGAACCGT ATTGTAAATC	1140
TCTGCCCTGCT CATCGCTAAC ATGCCCTAGA TAGATAGTCC GGGTCATATC ACTGACATAG	1200
TGGTCATAGA GACAGCCGAA GTCCATGGTG ATGGCTTCTC CCAACTCCAC TGGTTTGTC	1260
ATTGGATGGG CATGGGTTT AGAAGAATTG ATACCGCTAG CTAGGATCGT ATCAAAAGAT	1320
AAGCCAGATG CTCCCAACTC ACGCATGCCG AAATCAAGGA AGTTGGCAAT CTCAATTCA	1380

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GTTTTTCCCGTG	GTTTGATAAA	GTCAAGCGCA	TCCCGGAAAG	CTTGGTCTGA	GATAGAACAA	1440
GCCTTGGCAA	TCGCTGCAAT	CTCTGCCTCA	TCCTTAATCA	TACGAAGACC	TTCCACAAAC	1500
TGAGTTTGTG	GAAGCAAGTT	CAAACCTGCA	AAAGCTGCCT	GCATACGGTG	GTAATAAGAC	1560
ACTGAAATCT	CATCTTCAAA	ACCGATACGA	GTCAAGGCCA	TGTCTTAAC	AATTCTGCA	1620
ATGACAGCCA	ATTCATCAGC	ATCAGCCACA	ATCTAAAC	CACTGGTTTC	TTGCTTAGCT	1680
GCGATGATAT	AGCGAGAGTC	TGTCACTAAG	ACCTGACGGT	CACGACTGAT	AAAGACTGTT	1740
CCGTTTGAGC	CCCAAAACCC	AGTCAAATAA	TAGACGTTT	TAAGATTGTT	GATGATGATA	1800
CCATCTAGCT	CTTTTCTTG	CATTTTAGCT	AGAAATGCTT	GTACGCTTT	ATTCATGATG	1860
TAACCTTCCT	TTCAAATAGT	GTCCTGTATA	GCTGGCTTCG	TTGGCAGCTA	CTTCTCTGG	1920
AGTTCCCTGTT	ACGATGATGG	TTCCACCACC	GACACCGCCC	TCAGGCCCCA	AGTCATGAT	1980
ATGGTCTGCC	GTCTTGATAA	CATCCAGATT	GTGCTCGATG	ACGAGGACTG	TATTGCCATC	2040
GTCTACAAAG	CGAGCTAAAA	CCTTGAGCAG	GCGAGCAATG	TCCTCTGTAT	GAAGCCCTGT	2100
CGTCGGCTCA	TCCAGAAATGT	AGAAAGATTT	TCTCTGCGAT	CGTTTGTGGA	GTTCGCTAGC	2160
TAACCTTCATA	CGTTGGGCTT	CTCCCCCAGA	AAGGGTGGTA	GCTGGCTGTC	CCAAGGTAC	2220
ATAGCCTAGC	CCTACATCCT	TGATGGTCTG	GAGTTTGCCT	TGAATTTTCG	GAATGTGTTG	2280
GAAAAATTCT	ACCGCATCGT	TGACCGTCAT	ATCCAAGACC	TGCGAAATAT	TCTTTCCCTT	2340
GTAGTGAACT	TCTAGGGTTT	CACTGTTATA	GCGGGTCCG	TGGCAAACCTT	CACAAGCCAC	2400
ATAAACATCT	GGCAAGAAGT	GCATCTCAAT	CTTGATAATC	CCGTACACCTG	AGCAAGCTTC	2460
ACAGCGACCT	CCCTTGACGT	TGAAACTGAA	GCGCCCTTC	TTGTAGCCTC	GAATCTTGGC	2520
TTCATTGTC	TGAGCAAAAA	GGTCACGTAT	ATCGTCAAA	ACTCCTGTAT	AGGTAGCTGG	2580
GTTAGACCTC	GGCGTCCGTC	CGATAGGGCT	CTGGTCAATA	TCAATCAAAC	GGTCGACATG	2640
CTCAATCCCT	GTAATAGTCT	TAAACTTACC	AGGTTTGTCT	GAATTACGGT	TGAGCTTCTG	2700
GGCAATGGCT	TTTTTGAGAA	TGCTGTTGAT	TAGAGTCGAT	TTCCCTGAAC	CCGACACACC	2760
TGTCACTGCG	ATAAATTTC	CTAGTGGAAA	GCGAGCCGTG	ACATTTGCA	AGTTGTTCTC	2820
ACCGCGCTCCT	ATCACCTCAA	TAAAACGACC	ATTCCGACA	CGGCGCTCTT	CTGGTACTGG	2880
GATGACACGT	TTGCTGACA	AGTACTGACC	TGTGATAGAC	TTGCTGTTGC	GAGCCACTTG	2940
CTTAGGTGTA	CCTGCTGCAA	CAATCTCACC	ACCAAAAACA	CCGGCACCGAG	GACCAACGTC	3000
AATCAGATAA	TCAGCCTCAC	GCATGGTATC	TTCGTCGTGT	TCCACCACGA	TAAGAGTATT	3060
GCCCAAGTCA	CGCATCTTTT	TCAGACTGGC	AATCAGGGCA	TCATTGTCCC	TCTGGTGAAG	3120

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ACCGATTGAC GGCTCGTCTA GGATATAGAG GACACCTGAT AGGTTGGAAC CAATCTGGT	3180
TGCCAACGA ATGCGCTGAC TTTCCCCACC TGAAAGGTT CCTGCTGAAC GTGACAGGGT	3240
TAGATAGTTA AGACCCACAT TATTAAGGAA GGTCAAACGA CCCTTGATTT CTTGAGAAAT	3300
GGGACGAGCA ATGATGGCTT CATTTCAGA CAAAGTTAAC TGGCTCACCA AGTCCAAGTG	3360
GTCAGCGATA GACAGGTCTG AGATTTCTCC AATATGTGGC CCTTGCTGGC CGCCCACACG	3420
GACAGACAAG GCCTGGTCAT TGAGACGATA GCCTTGACAG GTTCCGCAG TCAGCTCATT	3480
CATGTAGAGA CGCATCTGAG TGCAGGTGTA ATCGCTATTG GTTTCATGGT AACGACGTTT	3540
GATATTATMG ATAACCCCT CAAACGGAAT GTCGATATCG CGCACGCCAC CAAATTCTATT	3600
CTCATAGTGG AAATGGAATT CCTTACCATC TGACCCATAG AGAATCAAGT TCTTATCTTC	3660
TTCTGACAGG TCCTCAAAAG GCTTATCCAT AGCCACTCCA AAGACTTCA TGGCTGCTC	3720
TAACATGTTT GGATAGTAGT TGGATGAGAT AGGATTCCAA GGTGCTAGCG CTCCCTCACG	3780
TAAGGTTTG CTAGCATCTG GCACTACCAA ATCAGTATCC ACCTCCAGCT TGATGCCAA	3840
GCCGTCACAC TCACTACAAG AGCCAAAAGG AGCATTGAAA GAAAAGAGAC GAGGCTCTAA	3900
CTCTGGGACA GTAAAACAC AACTGGACA GGCATAATGC TCAGAGAAC ACAACTCCGA	3960
GTCGTCCATG GTGTCGATAA TGACATAACC TTCTGCAATA CGAAGGGAG CCTCAATGGA	4020
ATCAAAGAGA CGACTACGAA TGCCCTCCTT GATAACAATA CGGTCAACCA CGACATCGAT	4080
ATTGTGTTGC TTGCTCTTAG ACAACTCTGG CACTTCGGTC ACATCATAGA CTTCCCCATC	4140
CACACGGACA CGAACATACC CGTCTTCTG AACCTTCTG ATAACACTCT TATGTTGGCC	4200
TTTTTCTTG CGGATGACAG GAGCCAAGAT CTGCAAGCGC TGGCGTTCAAGTCAA	4260
AACCTTATCA ACGATTTGCT CCACAGAAGA AGCATTGATA CCTCCATGTC CGTTGATACA	4320
GTAAGGCAGTC CCCACACGTG CGTAGAGGGAG ACGCAGATAG TCATTGATTT CAGTCGTCGT	4380
TCCCCACCGTC GAGCGAGGAT TTTTACTAGT CCTTTTCTGG TCGATGGAA TAGCTGGCT	4440
GAGACCATCA ATGGCATCTA CATCTGGTT TTCCATATTT CCCAAGAACT GACGAGCGTA	4500
GGCGGACAAA CTCTCTACAT AGCGACGTTG TCCCTCCGCA TAGAGAGTAT CAAAAGCCAG	4560
ACTGGACTTC CCTGAACCTG ACAAGCCAGT CACGACAACC AACTTGCTC GCGGAATCTC	4620
CACATCAATA TTTTTAAAT TATGGGCACG CGCCCCATGA ATGACAATTT TATCTTGCAT	4680
CTTTGTTCTT TCTAGTCCAT TATTGCTTAC CATTATACCA AAAAAGTGA GATTCTATTA	4740
CCCAAAAGGC CGATTTGTA GTATAATAGT ACAGTGTGAA AAAATCTGAA AAATGAGAAA	4800
GGATAAGGAA TATGAAACAA GTTTTCTCT CTACAACAC TGAATTAAA GAGATCGATA	4860
CGCTTGAACC CGGTACTTGG ATCAATCTCG TCAATCCGAC TCAAAATGAA TCACTCGAAA	4920

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TCGCCAACAC CTTCGATATT GATATTGCTG ACCTTCGAGC ACCGCTCGAT GCGGAAGAAA	4980
TGTCTCGTAT TACCATTGAA GACCGAGTATA CCCTGATTAT CGTAGACGTG CCGGTACGG	5040
AGGAAAGAAA TAACCGCAC CACTACGTAA CCATCCCGCT TGGTATTATC ATCACTGAGG	5100
AAACCATTAT CACTACGTGT TTGGAACAC TACCTGTCCT TGATGTCTTT ATCAACCGTC	5160
GATTGCGTAA TTTCTATACC TTCATGCGTT CACGTTTAT CTTTCAAATT CTTTATCGCA	5220
ATGCAGAGCT TTACCTAACCA GCCCTTCGTT CAATCGACCG CAAGAGTGAA CAAATCGAAA	5280
GTCAACTGCA TCAATCAACT CGTAATGAAG AATTGATTGA GCTCATGGAA TTGGAAAAAA	5340
CTATCGTCTA TTTCAAGGCC TCCCTCAAAA CAAATGAGCG CGTGATTAAG AAATTGACCA	5400
GTTCAACCAG CAATATCAAG AAATACCTTG AGGACGAAGA CCTGCTGAA GACACCCCTGA	5460
TTGAAACCCA ACAGGCCATC GAGATGGCAG ATATTTATGG AAACGTCTTG CATTCTATGA	5520
CAGAGACCTT TGCCCTCTATC ATTTCTAACCA ACCAGAACAA CATCATGAA ACCTTGGCCC	5580
TTGTGACCAT CGTCATGTCC ATCCCAACCA TGGTCTTTTC TGCCCTACGGG ATGAACCTTA	5640
AGGATAATGA AATCCCCCTA AACCGGAGAGC CAAATGCCTT CTGGTTAACTC GTCTTTATCG	5700
CCTTTGCTAT GAGTGTCTCG CTCACTCTCT ATCTCATCCA TAAAAAATGG TTCTAAGAGG	5760
AGTTCCATG TCTCAAATTG ATCTACAAAA ATTAACTAAG AAAAACCAAG AGTTTGTCGA	5820
CATTGCTACC CAACAATTCA TCAAAGATGG GAAAACAGAC GCTGAAATCC AGACTATTT	5880
TGAGGAAGTC ATTCCCCAAA TCCCTGAGGA GCAATCTAAA GGTACAACGT CCCGTTCCCT	5940
ATACGGCGCA CCAACTCATT GGGCTCATAG CTTCACTGTC AAAGAGCAGT ACGAAAAAGA	6000
GCATCCAAAA GAAAATGATG ACCAAAAACT GATGATTATG GACTCAGCTC TTTTCATCAC	6060
TAGCCTCTTT GCCCTTGTCA GCGCCCTCAC AACCTTCTTT GCGGCAGACC AAGCTTTCGG	6120
CTATGGATTG ATTACTCTTC TATTAGTTGG ACTGGTTGGT GGATTTGCCT TCTACTTGAT	6180
GTACTACTTT GTTTACCAAT ACTATGGACC AGATATGGAT CGCAGTCAAC GTCCACCTTT	6240
CTGGAAATCT GTACTAGTTA TCCTAGCTTC TATGTTCTT TGTTGCTTG TCTTCTTGC	6300
AACAAGCTTC CTACCAAGCTA GCCTTAACCC AGTACTGGAT CCATTGCCAC TAGCTATTAT	6360
TGGAGCAGCC CTCCCTAGCCC TTGCGCTCTA TCTCAAGAAA CGCTTGAATA TCCGTAGTGC	6420
AAGTGCAGGA CCAACACGCT ATCAAGAATA AGAAAACGAT AAAAGCAACT GCAGGGCGG	6480
TTGCTTTTC ACTTACTTTT TTGAGTTATA TTCATGAAA ATCAAAGAGC AACTAGGAA	6540
GCTAGCTGCA GGTTGCTCAA AGCACAGCTT TGAGGTTGCA GATAAAACTG ACGTGGTTG	6600
AAGAGATTTT CGAAGAGTATT TAAAAGTATT CTTCTGAAAT CCCACATAGC TTTCTCTTAT	6660

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ATTTTGTGAT AAAATAGGCT CAATCTATTT CTAGGAGGAT GAGATATGGT TTCTACTATT	6720
GGTATTGTTA GTTTATCTAG TGGCATTATC GGAGAGGATT TTGTCAAACA CGAAGTGGAC	6780
TTGGGTATCC AACGCTCTCAA GGATCTGGGA CTCAAATCCC A TCTTTTT	6827

(2) INFORMATION FOR SEQ ID NO: 61:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 11864 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 61:

CTGGCTAGTT GCATAGAGCA AAGTTGCTTC TTCATCAACA AAACCGTTCA TTTCAAAATA	60
GGAAAGCAGC TCATCAGGAC TCTCCAAACG AATCCCTTG TAATCCAGCT CAACTGCCAC	120
CTCTTCAAG GCTGCAAGAA GAAGTGTTC CAGGCCCTGT CTCTGATGGT CAAACTCGAT	180
GACTAAAGAA TGTACTTTA GACATPGCGG ATTGTCTGAC TGGGGACTTG ATAAAATATA	240
GCCTAAAAAGT TGATTTCAT CCCTAGCTAG AAGAAAGGTA TCCGCACACT TACGGATACT	300
TTCTTCTAAA ATATGGAAA GTTGCTGCTT TTCAGCTGGA AAAGACGAGG TCTGAAGTGC	360
CCCTATCTCA GGCAAATCG ACTTGCTTGC CTGAATGATC TTAATTGGAA TTTCCATGGG	420
AACATCCTAT TGAACATTGC TTGTCAGTT AGACAAGAGA CGCTCAAATG AGTATTCTATA	480
GGTTTGGATG TCTCCTGCTC CCATAAAAGAC GTAAACAGCA TTGTCATGGT CTAGGAGTGG	540
AGAAAACATT TCAACAGTAA TCACTTGGTG TTTTTGTTG ATTTTGTGG CTAGGTCTTC	600
TACCTTAACG TCACCCTGAT CTACTTCACG AGCCGAGCCA TAAATTGCG CTAGATAAAC	660
AGCATCTGCT TGGTTAAAG CATGGGAAA GTCGTCCAAC AAGGCAATGG TTCTTGTAAA	720
GGTATGCGGT TGAAGAGCT CTACAATTTC CTTGCTTGGG TATTTCAGAC GAGCCGCATC	780
CAAGGTCGCA ATAATTTCTG TTGGATGGTG GGCAAAGTCA TCGATAATCA CTGTATCATT	840
GACAATTTC TCAGTGAAAC GACGTTAAC ACCGGCAAT GTTTCAAGT GCTCACGCAC	900
CAAGTCAAA TCAAATCTG CTGTGTAAG AAGACCAATA ACGGCTGTG CATTCTGAT	960
ATTGTGACGA CCAAAGGTTG GAATGTGGAA TTGCCCCAAG TTTTGTCCAC GGAAATGAAC	1020
GGTGAAGGTT GAACCAAGTTA TTGAAACGAAG AAGATCACTA GCTACAAAGT CATTGCTTC	1080
AGCTTCAAAA CCATAATAAT AAATTGGTGC ATCAGACGTA ATCTTACGCA ATTCAAGCATC	1140
TTCACCATAG ACAAAAAGAC CCTTGGTGAT TTGTTGGCA TAGTCGTTAA AGGCATTAAA	1200
AACATCCTCG AGACTTGTGA AATAATCTGG ATGGTCAAAG TCAATGTGG TGATAATAGA	1260

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GTATTCTGGG TGGTAAGGCA TGAAGTGACG CTCATATTG TCAGATTCAA AGACAAAATA	1320
TTTGGCATTG GCCGAACAC GACCTGTCCC ATCTCCAATC AAGAAGCTGG TATCTGTAAT	1380
GTGAGACAAG ACATGAGACA ACATACCTGT CGTTGAAGTT TTTCCATGTG CTCCTGCTAC	1440
TCCCCATGCTA ACAAAAGTCAC GCATAAAAGCT ACCTAGAAC TCATGGTAAC GTTGTAGCT	1500
GATACCATTG TGGTCCGCAT AGGCAATTTC GACGTTGTTA TCTGGACGAA AGGCATTCC	1560
AGCGATAATT TCCATATCAC CGCTAGATT TTTTCATCA AAAGGAAGAA TGGTAATTCC	1620
TGCCTGCTCA AGACCGCGTT GGGTAAAGTA GTACTTTCA ACATCTGATC CCTGAACCTT	1680
GTGCCCATC TGTTGCAACA TCAAGGCCA GGCACTCATC CCTGATCCCT TAATTCGAT	1740
AAAATGATAT GTCTTGACA TGTTTCTCC CCTATTCTGT CATTCTGGTC AGATTCAACT	1800
CTTGGGCAAC CCGACGTTCT TGTTCTGTTT GTTTACTTTT TTTATTGTAG ATTTGGCTCT	1860
TCTTTAGAAA ATCATAATTG TTTTCCTTG GAGCAGGTGC TGACACTTCT TCATTCTTGG	1920
TAGGGATAGA ATGAACCTCT TCCGCCAAGA TATAATGAGA CTGGGTCAAT TTTTGCTAT	1980
ATTTGACAAA TTCAACCAGGA TTTCCTTTT GGAAAGGAGC TGTCGGTTGA TTGCCCTGTC	2040
TAACTAGACT GGGCTGAGAA TGACGTCTCG CAAGGCTGAA ATCCTGAGTT AGGTAGTTAG	2100
CAGAGCGTTT CTPTTCAAG TCCGCACCGG CCTCTTCACG CGCACCCCTCC GCATAGCTCT	2160
TTCCTCTTTT TTAAACCCCT AAAGGAGCCT TTTTAGGTTT TTCGACTTGC TTTCATCG	2220
GTTTTACTGG TTTCCTTCA GCAATAGGAG CCCATTCTAA ATAATTTTA TCTCGATACT	2280
CACCCCTTGAT ATTACTGATC AGATCAGACT CATCATAGAG ATTCATGACT GGCATTTCA	2340
TCAACATGAC CTCGTCATCT GACACCAATG GAAATCGTTC TTGTTTCATT TTCTATTCC	2400
TTTCAACACT TCATTATAGC GTATTGTCTT GATTTTCAA GTGCTGGCTT CAGAAATTCC	2460
CAAAATTTCT CTAATTTCTG CTAGGGTCAG ACTACCACGT GACTCTGTGC CGTCCAATAC	2520
TTGTGACACC AGATGTTCTT TTTGTTCTTG GAGTTCCCTGA ATTTTTCTT CAATGGTTCC	2580
CTTGGTCACC AAGCGATAGA CCTCAACCGT TTCTTCCTGA CCCATCCGAT GGGCACGGCC	2640
AATGGCTTGC GCTTCCACCG CAGGATTCCA CCAAAGGTCA ACCAAGATCA CTGTATCTGC	2700
ACCTGTCAGG TTCAAGACCGA CCCACCAGC CTTGAGGGAA ATCAGAAAGG CATCTCTTC	2760
TCCCTGGTTA AAGGCCCTTG TCATGTCTTG TCTTTCCCTG GCTGGGGTTG AACCCGTAAT	2820
TTTAAAGGAA GTCAGGCCA AGTCTGGCAG TTCTTGTCAT ATTTTTCTCA ACATTCCCTT	2880
GAACTGAGAG AAAATCAAGA CACGGTGTCC GCCGTCTGCC ACCTGTACCA GTAGGTCTCG	2940
GAGACTATCT AGTTGCCGC TGGCTCCCTG ATAATCTTCC ATAACACAGGG CAGGAGTGTC	3000

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ACATATTTGA CGCAAGCGCA TCAAACCAGA TAAAATTCC ACACGACTTC GCTGAAATC	3060
CTGTTCTGAC ACTTGAGCCA GATGGTCTCG CATCTGTTGT AACTGGGCAA GGTAAATAGC	3120
CTTTTGCAGG TCTTCCAGTT CATTTTTATA AACCAACCTCA ATCAAGTCTG GCAATTCACT	3180
CAGAACCTCT TCTTCTTGC GTCGCATCAC GAAAGGCTTG ATAAACTGAG CCACTCGCTC	3240
TGCTGGCAAT TTCATAAATT CTTTCTTGCT TGGCAAAAGT CCAGGCATGA CGATTTGGAA	3300
AATAGACCAC AACTCACCCA GATGGTTTTC AATCGGAGTT CCTGACAAGG CAAAGACCGA	3360
CGGCACCACA AATTGTCTCA AGGTCTGGGC AATCTTGGTC TGGGCATTTT TCATGACCTG	3420
AGCCTCATCT AAGAAAAGGA AGTCAAAGGC CATCCCTGTA TAAACACTCAC TGCCCTGACG	3480
GAAGGTGGCA TAGCTAGTCA CATAGATTTG ATGGCTCTCG GCAAGAATCT CCTCACGACT	3540
TGCTTCAAA CCATGAACAA CAGTCACATC CAACTGTGGA GCAAATTCT GAAACTCATC	3600
TGCCCACTTG TAAATCAAAC CCGACGGAGC GAGAATCAAAC ACCCGACTTT CTTTGTAC	3660
TTGACTAGTC AAAAAGCAA TGGTCTGAAG GGTTTCCCAGTCCATAT CATCAGCCAA	3720
AATCCACCA AAACCATATAAT GATGGAGCAT CTGCAACCAG CCAATTCCCT TTTCTGATA	3780
ATCTCGCAAG TCAGCCTTGA CCTGAGTTGC TTGCAAAGGA AAGTCCTCTG GATGCGTCAA	3840
ATCCTGGGCC AGATTCTGGA ATTCTTGTGA AAAAGAAACA CGGTCTCGCC CTTCAAAGAG	3900
ATGAGCTAA CTGTAGGCCA AGGATTTCCG AGCCTGCAAG GTCCCATCTT TTAATTCAA	3960
TTGCCCCAGT TCCTGTAGAT TTTGGCGAAT TTTCTTGGTT TCTTCATCGA AAAAGTAAAC	4020
TTGATTAGAC GAATCAATAT AAAAATCCTG ATTGGCAACC AAGGCCTGCA TGGCTTGGTC	4080
GATTTCTCC TGGACAATAT TTTGAAAATC AACTGGATT TCCAAGAGAC CTCCCTTGGAA	4140
GGCAATCTGC ACCTGAGGAC TCGCTAGGCT ATAAAGCTCT TCTAGTTTAT CTGATAGGTC	4200
AACATGCCCG AGTTTTCAA AGACTGGAAT GATATCATGA AAAAATGAT AGACAGACTC	4260
CGCTTTAAG GCCTGACGCC AAGATTGAAA ATCGGCCCTCA AAGCCCGCAG CCAAACAGAC	4320
TTGGAAAAATT CTTTCTTCTA AGTCTGCGTC ACTTGAAAAG GGTAATTCTT CTAGCTCTTG	4380
TCGGCTAGAT ACCTGTCTAT TTCCATAATC AACTGAAATT TCTAAACGAA TCCGATTATC	4440
TTCTTCCCTG TCAAAGTAAA AAGAGGGCGC AAAAGTTTG ATTTGTAGAC GTTCTGGAGC	4500
TGAAACGGTG CCCATCTGGA TAAAAAGAGT CAGACAGGAG GCCAATTGTGCT CGATCACT	4560
GCTATCAAAT TGCAGGTATT TCTTCCCTG TTGACCCACA GGTAACGCTT TAATTTCTT	4620
GAGAAGACGC ATCTGCTGGT CTGTTAAAAA ATAAACCTGA CCTTTATGGAA AAAGTACTGC	4680
TCCCTGATAA AAGACATTGA CCCTAGGACT CTCACTGATT TCCATTCAA AATAATCCGA	4740
GTATTCTGTT ACTGTAAAGG CAAATAGATT GGCATCAGCA TGCATATCCT GAAAAGCAG	4800

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GGTTTGGTAG CTATCCACTT GATGGTCAA TTGAAAATGG GGCAAGGCCA TCAGTAAATT	4860
CACACCTGC TCAAAAAAGG TCAGAGGGAA AAAGAGGTGC CGACCTTGTT TTTGGAAAAA	4920
GAGGTCTGGA ACCAGCCCTT CCTCCGTTAG TCCGTGCAAG AAAGTCAAA GTTCTTGCT	4980
GGCATCATCA AAGGCTTCCC AAGAAAGAGA CTCCCTCATAA ATCTTGCCAA TCATATACGA	5040
CTTTCTCTGC TCGACAATCC TTAAAAAAAAG TGGAATATCC CGAATGACAT AGTATTTTG	5100
GCTATTGATT TGGCCGATTC TCAGAGTCCA CAAGATATGA TTGGTTCTG CTTCCACCTG	5160
ACCCACAGCT GATAACTCAT AGGCCATTC TGATTTGGA GATAAAATTC GATCCAAAAA	5220
CTTGCCACCC AAGGTCACCT TGGTTCAAC AGCCTTTTT TCTTCATGAC CTTCTTCCAG	5280
ACTCCACAAG ATTTCTGAC CACCGTCATC ATTTTCAGA AAATGCTCTA GCGCTGCCAA	5340
ATGCCACACAG TAGCCCCCTCT TTGAAAAAAATC ACAGAGGCA CAAAAAACCA AATCATCCTC	5400
TAAACTATAG CGCAGTTCTT CTTCTGCAAC GCGAGCGTAG AGCCGATTGT TCTTTTCCTT	5460
GATGATATCA ACCTTACCAAG TTTCATAAAAG GGCACACACT TCGATACGAA TTTTCCCCGG	5520
AATCAATTAA GCCATATTTT CACCTTTACC TTATTTTTT ATTATACCAT ATTTTCGCTC	5580
ATGAAAATAG CCTTCTAGGA AGACTTTCT CCTAGAAGGC TGGATTTTA ACGTTGGCA	5640
AAAGTAGCCA CAATCCGCTG ACAGACTTCT TGCAACAGAG ATTTGGCAT AGCTATATTG	5700
ATGCCGGCAT GGAGACTTCC TTCCCTCTCCA AAATCCAAAC CACGTTGAG GATAACCTTG	5760
GCTTCATTTTC TCAACAACCTC TTGCAATGTT TCATCAGTC GGTCTAAGC TGAAAAGTCA	5820
AGCCAATCA AGTAGGTACC TTGCGGTTTC ATGACCTTGA TTTTAGTCTC TTTTCCAAAT	5880
AGATCCATCA CATAATTGAT GTGGCTTCA AAAGACTTGCT TGAGTTCTC TAGCCAATCT	5940
TTACCGTATC GATAGGCAGC TTCTGTCGCC AAATAACCCAA AGCCTGAAAT TTCTGCTGA	6000
TTATTGGCCA ACAGCGTTT CTGAAAGCC AGTCTCAACT TAGGATTTTC AATGACTGCA	6060
TAGGAATTAAAT TTGTTCCAGC AATATTAAAT GTTTAGTGG CACTGCTCAA GACGATAGCA	6120
AAATTTTGA AGGCAGGATT GATGGTATTG AAAGACTGGT GTTGTGACC AAAGAGGGTC	6180
AAATCTTGGT GAATCTCATC CGAAACTAAC AAAACACCGT GTTTTGGCA GAGTTGGCA	6240
ATCTTCTCCA ACACCTCTTT TTCCCAAACA CGTCCACCAAG GATTGTGAGG GTTGCAAAGA	6300
ACATAGAGTT TAACCTCCTC TTCCACCAAAC TCCCTTTCAA GTTGGTCAA GTCAATCTCA	6360
AACAGACTAT CCTTTCCAC TAAGGAATTA GTAATCAATC TACGATTATT CAACTGACA	6420
CTGCGAGCAA AGGGTGGTAA GACAGGGGTG TTAATTAAAA CCGCCTCGCC TTCTTTGTA	6480
AAGGTTTGAA TAGCTGTTGA GATGGCTGGT ACCACACCCCT CGATAAAAGAC AAGAGCCTCT	6540

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TTGTCAAAGT	TGTAACCGTA	TTGTGTAGCT	TCCCACTTTT	GAACCTCCTT	AATTAAGTCT	6600
TCACTGGCAT	AGGTATAACC	ATAAACCAAGT	TGGTCTGCCT	AAGTTGCAC	GGCTTGGCGG	6660
ATTCAGGCA	AGACACACAA	GTCATATCC	GCTATCCAAG	CTGGTAGAAC	TTCACTATCC	6720
GTTTCTGTT	CTTCCATT	ATAGGTATGG	TGCCCTAAC	GGTTGGGCAG	GCTTGTA	6780
TCATATTTTC	CCATCTTGT	CTTACCTTC	TATGGCTTG	CGCAAATCTG	CAATCAAATC	6840
TCTAGCATTCC	TCAATCCAA	TAGACAAACG	CAAGAGGTCA	TCTGTCAAC	CATAAGAATG	6900
GCGTACCTCT	GCTGGAAAT	CAGCATGAGT	TTGAGTCGTT	GGATAAGTAA	TAAGACTTTC	6960
CACTCCACCC	AAACTTCCG	AAAAGAGAA	GACCTTGAGA	CTGTTCAAAA	TATGAGGAAT	7020
GCGTGTTC	TCGGTACTT	TAAAGGAAAT	CATGCCCTCA	CGACCAGTGT	AGAGAACTTC	7080
CTTAAC TGCT	GGAGAACCT	TCAAAAGGC	AACCACCTCT	TGGGCGTTAG	CTGTTGAGCG	7140
CTCCATACGA	AGAGACAAGG	TCTTGAGACC	ACGAAGAAC	TGGTAGCTGT	CAAATGGAGA	7200
CAAGACTGCC	CCTGTTGTAT	TAAGATTGTA	AAAAAGCTTC	TCGTATAGTT	CTAAACTATT	7260
GGTCACAACC	ACTCCAGCCA	AGACATCATT	GTGGCCTGCT	AGATACTTGG	TTGCTGAATG	7320
GAGAACGATA	TCTGCCAT	CTTCATCGG	ACGTTGGTAG	ATAGGGCTAT	AGAAGGTATT	7380
GTCCACCAACC	ACTTTGGCAC	CCTTAGCATG	AGCCAATT	GCTAGTTTT	CGATATCAA	7440
TTCCAACATC	AAGGGATTGG	TTGGGTTTC	GATATAGAGA	ACATCCACAT	CCTTTCTAA	7500
CTCGGCAATC	AACTCTTCTT	CTGTATTGGC	ATAGTAAAA	TGGAAATGAC	CTTCCTGCTC	7560
CACTTGGTTA	AACCAGCGAA	AAGAACCAACC	GTAAAGATCA	CGCACTGCCA	AGACCTTACT	7620
TCCTACTGGA	AAGACGCTAA	AGGCCAGTAC	AATAGCTGAC	ATCCCTGAGC	TAGTCGCTAG	7680
GGCATAGTCT	GCTGACTCAA	TAGCCGCCAA	GA CTTCTCA	GCCTTACTAC	GAGTTGGATT	7740
TTTAGTGC	GTATAGTC	ACCCAGTAGA	TCGACCAAC	TCTGGATGCT	GATAGGTCGT	7800
TGAAAATGA	AGTGGTGTCA	CCAAAGCACC	TGGTGCCTCA	TCAGACTTGA	TCCCTGTTG	7860
TGCTAAAATT	GTGTTAATGT	GTAATTCTT	GCTCATACAA	TTCCTCCAAA	TCTATAGTAA	7920
CTATTGTACC	ACTTATTTTG	TATCCTTCGT	TTTCTGTTT	TCAAGAGCTA	GTTATAGTT	7980
CAAAC TATAT	AAAAGGGAG	TTTTCTCTGC	TCCCTTTAAT	AGACTATAAA	ATGGTGAATC	8040
TCAAAAGACA	CCTTCACTCT	ATCATTTGCT	CCTGCACAA	ACGAGCATAA	CGCTCATGAT	8100
TTTCCAGTAG	TTCCCTATGA	GTTCCCTGAGC	CAGTGTATTT	CCCCTCTCT	AAGAAGAAA	8160
TACAATCCAC	ATCTTTTAC	GTTGACAAAC	GATGCCTAT	AATCACAACC	GTCTTCTCCT	8220
TTAGTACAGA	ATAGAGGCTA	CTGATAATCG	CATACTCAGA	ATCCGCATCA	AGATTAGCAG	8280
TGGCTTCATC	AAATATAAGA	ATTCAGCAT	CTTTTAAGTA	GGCTCTAGCT	ATTTGAAGTC	8340

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TTTCGTTCGC	CCCCCTGACA	AGAGTCGTCC	GCGTTCACCA	ACTTCAGTAT	CTAGTCCCTC	8400
TTTCATGGAG	CGAATCTCAT	CACCTAGTGA	TACTAAGTCT	AGCACTTTCA	TCAATTCTAC	8460
ATCAGTTACT	AAGCGATTCA	AACCAGAGACA	AAGATTGTCA	CGAATACTGC	CAGATAAGAC	8520
TGCATTATTT	TGTGAAACCC	AAGCGATTTT	ACTTCTCCAT	TCTTTTAAGT	TAAAATCATA	8580
TATACTTGAT	TGCTCCATTA	GAATATCTCC	TGAAAGCGGT	TTATAAAACC	GCTCTAACAA	8640
ACGCACAAATC	GTTGATTTTC	CTGATCCAGA	TGGTCCAACA	AAAGCAATT	TTTGCCCCTT	8700
GAAAATTGAA	CAAGTAATAT	CCTTTAAGAC	AGGTCGATTT	TCATCATAAC	CAAATAGAC	8760
ATGGTTAAAA	TTCAACCCCTC	GTCTGATAC	CGATTTCCCT	CCCTCAAATT	TTTCTTTAGG	8820
AACTGCAAGC	AAGTTCTCCA	GTGCAACTGA	AGATCCCTTG	CTCCTAGAAT	AAACAGTTAC	8880
AAAATTAGCT	ATATTACTAA	TAGGATTAAG	TAATTGAAAG	AGGTAAATCA	AAAACGAAAC	8940
CAAGGTTCCC	ACAGATATAT	ATCCTGCGCT	GACCCGATAA	CCCCCATAGG	TTAGCATCAC	9000
AGCTATAGTC	GCAAAGATAA	ATAAGAGAGC	AAACGGGGTC	TCAAAAGAAG	TAACCCCTATC	9060
TGATTTCACT	GAATTGTTTT	GTACCCCTTC	AATACAATTA	TCCAAAACAT	CCTGTACACT	9120
TTTCTCTGCT	TGGTTAGTCT	TAATTAATTC	ATGTTCTTGA	ATCTTTTCAG	TCAATTGCC	9180
TGTTAAATT	CCTCCTGTAA	ACGACGACTA	TACTTTTCAC	TGATATTGGA	AAGGGGCAAG	9240
ATAAATAAAC	TCATACAAGG	AAGACTGATG	ATAAAAGTA	GAGAAAGATT	CCAATCAAGA	9300
CTAAATAAGA	CTACANTCGA	ACCAAGTACC	ATAACTAAAC	TCAGAATAAT	ATTTGGGAAA	9360
GTCGTAATTA	AAAACTCAGG	AATGACACTC	GTGTCATTGA	CAATGGCAGA	AGTCAACTCC	9420
CCACTTTGGC	TCTTATCAA	GAAGGATTTC	TCTACATAAA	TCAACCCCTC	TATCACTTT	9480
TTCCCTGATTT	TTGCTATCTT	TTTTTCACCC	GATTGACTAA	ACAGATAGTA	ACCAATAGAA	9540
GAAAACAAGG	CTTGACCAAT	AAAAATCAA	AACGATTGAA	ATACTTTGGA	GCCTATATT	9600
TCAATAGAAC	TCCCACCTAT	TAATCCTTT	AAGATAAGGG	GAAGCAACAA	AGCAAGTAGA	9660
CTAGACAGAA	CAAGTAAGAA	ACTCCCCATA	ATCACCTTAG	TATCTACTCT	TAATAATT	9720
AATTTCATAA	ATACTCCTTA	TAATATTTC	ACGGATAAAG	TCGGAATAA	CTCAATTG	9780
GGATAAAATC	TAATAAAATCT	TCCTATAACA	AAACGCATAA	CATCTAGGAT	TTTATATACC	9840
TGATATTATG	CGTTTTAAG	CACAAAGACT	TCTTACACAA	ACTTATCTAC	AATTAGATT	9900
TATTTGACAT	GTGTTGCCAA	TTCTTCTTGG	GCTTTTTAT	TGGATTCTTC	TTTTCTTTC	9960
AACCATTTT	CTCTGGCTTT	TGCATATTG	TCTGTTGTGA	CAATCTTATC	TTGTACTTTG	10020
AGGTATTAT	ATGATTCAAC	CCCTTTGTA	CCGGTAAAC	CATAGGCAGC	AGCAAATGGT	10080

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ACGGTTCTTC TCAATGATGG TGTTCCCCA CGCGAAACAC TTGGAAGAAC TAAAGAACTA	10140
TCAATCAACC AAGCTTGAAT ATCAGCATAT TTCTCATAAC GTTGGCCGG ATCTTGCTCT	10200
TTATTAGCTT CTTCAACAT TTGAGTATAG ACATCCAGTC CAACTGCCCT AGCCTTGTCA	10260
TTGGCCCTCAC CAGGCCTCTAG TCCAAGATTT TGCAGAAATC CTCCACTATT AGTATTAAAA	10320
ATATCGAGAT AGGTTGACGG GTCTTGATAA TCAGGTCCCC AACCGCCATG ATATAAATCA	10380
TAATCTTCT GAGCAGCTGT TTGAGCAAAG TAGCCTGAAC TGTCAAACTC ATCTGATGTT	10440
AATTGCTGAA TGTCAATCAC TACATTATCA GAACCTAAAA CAGATTCAAT TGATTGTTG	10500
ATAGAACTAA CTCCCTGTAT GCCTACTTTA TCTGTTACTT CCACAGTCTT ATCCAAGTGG	10560
ATGGGAATT GAACACCCCTT TGCTTCGAGT TCTTTCTTAG CTTCCGCAA CTTAGCCTTG	10620
GCTTTCTCAAG GATTGCTAGA AGGGCTCTGA CCATCCGCAA AGTTGATACC TTGCCATTCC	10680
TTACCATAGT TGACCATCTT AGAGGCTACA ACTTCACCAA AGTCTTTCC CTTGATACTG	10740
ACAAAGTTG GAGGAACCAC TAGGTTACGC AAAATCTTTG TTGACACCTTC TTTCCCTTC	10800
GACTGAGCCC CATAAGATGT TCTGTCAAAA GCAAAATTGA TAGCCTGACG GAAGTTTTA	10860
TTGAGAACTG CTTCCCTGAGT CGATTTCTTT TCAATGTCAC TTGTTTCTAG AGTATAATTG	10920
TAAGACTTCC TATCTAGGTT AAAATTAAAG AAATATGAAG TTGAATTTTG CATACTATAG	10980
ATGATATTGT TTTTGTATTT TTCTTTAATC CCTTCATAGC TGGAGCTGTT AGGAAAAAGA	11040
CGAGCCGTAG TATAAGCACC AGCTGTAAAA TTACGTTCCA GTGATTCTTG GTCGCTACCA	11100
TCATAGTAGG TCAATTCAC ATCGTCTACA AAGACATTCT TAGCATCCCA GTAATTAGGG	11160
TTTTCTTAT ATTCAATAGC AGATTTGAG ACAAGTGCTT TCATCAAGAA AGGTCCATTG	11220
TACAAAATAC TAGATGGATC CGCCTTCCCA AAATCATCCC CTTTGATTT CAGGAAATCT	11280
GCATTAACAG GAAAAGTAT CGTTGCAAGT GTTTTGAAT TCCAGTAAAG TTCTGGTTA	11340
ACCAAAGTAT ATTGAACCGT TTGGTCATCA AGTGCCCTGA CACCGACAGT TGAAAAGTCG	11400
CTTGTCTTAC CAGTGTATAA GTCATCCAA CCAGCAACAG AGTCCTGCAC TAGATACAAG	11460
GCTTCTGATT TTTTATCAGC TGCATATTGC AAACCTGTCA CAAAATCCG GGCAAGTTACA	11520
GGCGCATATT CTTCTCCCTC AGAAAGTAAAC CACTTGGCAT CCTTACGAAG TTTGTAGGTA	11580
TAGGTCAAAC CGTCTTGAGA AACAGTCCAA TCCCTCTGCTA ATGATGGAAT AATATTCCCA	11640
TATGGTCAT TTTCTAATAA CCCGTCTACC AAATTTGCAA CAATATCGGA TGTTGCTGCG	11700
CGGTTTCTG CTAGATAGTT CAAGCTAGAT GGATCACTTG AATAAACATA GTTGTAGGTT	11760
TTTGACGCCG TGCTAGAATT TCCACACGCCG CTCATAAAA CTCCTGTACC CAGGACAAGA	11820
CCTGCCAAGG TTAGATATTT GCTCTTAGAC TTTTCATTT CCGG	11864

(2) INFORMATION FOR SEQ ID NO: 62:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 2412 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 62:

TAAC TGC ACT AACATAATA TAAGGAGAGA AAATGTCTGC AATAGAACGT ATTACAAAAG	60
CTGCTCACTT AATTGATATG AACGATATTA TCCGTGAAGG GAATCCTACT CTACGCCGA	120
TTGCTGAGGA AGTCACTTTC CCCCTATCTG ACCAGGAAAT CATCTAGGC GAAAAGATGA	180
TGCAATTCCCT TAAACATTCC CAAGATCCTG TCATGGCTGA AAAAATGGGA CTCCGCGGTG	240
GTGTTGGACT GGCTGCTCCC CAGTTAGATA TCTCAAACG CATTATCGCT GTTTTGGTAC	300
CTAATATTGT TGAAGAAGGC GAAACTCCAC AGGAAGCCTA CGATTGGAA GCCATTATGT	360
ACAATCCAAA AATCGTCTCT CACTCTGTT AAGATGCTGC TCTTGGCGAA GGAGAAGGTT	420
GCCTGTCGT TGACCGTAAC GTGCCTGGCT ATGTTGTTCG CCATGCCCGC GTTACTGTTG	480
ACTACTTGA CAAAGATGGA GAAAAACACC GTATCAAAC CAAAGGCTAC AACTCCATTG	540
TTGTTCAAGCA TGAATTGAC CACATTAACG GTATCATGTT TTACGATCGC ATCAATGAA	600
AAGACCCATT TGCAGTTAAA GATGGTTTAC TGATTCTTGA ATAAAGAAAA TCCCGTTGCA	660
AGACGGGGTT TTGTGTTATA ATAGAGGCAT GAAAACAAT GATATTGTCT ATGGTGTCCA	720
CGCCGTTACC GAAGCCCTCC TTGCAAATAC AGGAAACAAA CTCTACCTCC AAGAAGATCT	780
CCGAGGTAAG AATGTTGAGA AAGTCAGGA ACTAGCTACA GAAAAGAAGG TGTCCATTTC	840
TTGGACATCA AAAAAATCTC TCTCTGAGAT TACTGAAGGT GCTGTTCATC AAGGTTTGT	900
TCTACGAGTG TCTGAATTG CCTATAGCGA GCTAGATTAC ATCCTTGCAA AAACACGCCA	960
AGAAGAAAAT CCACCTCTAT TGATTCTAGA TGGTCTAACCC GATCCCCATA ATCTGGGTMTC	1020
TATCTGCGA ACAGCCGATG CGACCAATGT TTCAGGTGTC ATCATTCCCA AGCACCGTAC	1080
TGTCGGAGTA ACTCCTGTCG TTGCCAAAAC AGCCACAGGT GCTATTGAAC ACGTtCCAAT	1140
TGCCCGAGTG ACCAACCTCA GTCAAACCTT AGGATAAACT TAAGGATGAA GGTTTCTGGA	1200
CCTTTGGAAC GGATATGAAC GGTACTCCTT GCCACAAGTG GAATACAAAA GGGAAAATCG	1260
CCCTCATCAT TGGAAATGAA GGAAAAGGTA TCTCTAGCAA CATCAAAAAA CAGGTCGATG	1320
AAATGATTAC CATTCCGATG AATGGACATG TTCAAAGCCT TAATGCCAGT GTTGCTGCGG	1380

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CCATTCTCAT GTACGAAGTT TTCCGAAATA GACTATAAAA AAGTTTCCAG TCATCTGATT	1440
GGAAACTTTT TTATGATTA CTATGTTCTG TAATGAATT ATAGGCTTCT TGACCACCGA	1500
TAGCTCCATC TCCAACCGCT GTGTGTTACTT GGCGAAGGTC TTTCAAGCGA ACATCTCCAA	1560
CTGCAAAGAT ACCGTCGACT GCAGTTTCA TGTGGTTATC TGTCACAATC CATCCTGCCT	1620
GATCTGGAT ATTCAATTCT TTAACAAAAT CGCTAAGAGG GTCCAAACCA ACATAGATAA	1680
AGACACCCACC GAAGGCTTGT TCTGTCACCT GACCTGTTT CACATTTCA AATACGACTG	1740
ATTCTACTCG GTTTTCACCC TTGATTTCCC TTACTACAGA ATCCCAGATA AAGCTGATT	1800
TTTCATTGCG AAAGGCGCGA TCTTGAAAA CCTTTGGGC ACGAAGTTGG TCACGACGGT	1860
GAACAATGGT AACAGTCTTA GCACAAACGAG TCAAGAAGAG GGCTTCTTCA ACAGCTGAAT	1920
CTCCACCCACC AACTACCAAT AAATCTTGGT CACGGAAAGAA AGCACCATCA CACACAGCAC	1980
ACTAAGAAC ACCACGACTG TTCAGTTCTT CTTCTCCAGG CACTCCAAA GGACGGTGT	2040
TAGAACCGAG TGCTACGATA ACTGTACGTG TTTCATATGT TTGGTCATCA GTCATCACTT	2100
TCTTAAAATC ACCATGGCTT CGACATTTTC AACATAACCA TAAATGTGCT CAACACCAAG	2160
ATTTTCAAGT GGTTCAAACA TCTTTCTCAGC CAATTCAAGGT CCACAAATAT TAGCGTATCC	2220
TGGGTAATTT TCGATATCAG ATGTATTATT CATCTGACCA CCTGGCAGAC CACCTTCAT	2280
CAAAGCTACT TTTAGATTGC TTGAGCAGC ATACAAGGCC GCAGTCATCC cTGCAGGTCC	2340
AGCACCGATA ATAATAGTAT CGTACATATA GATTCCCTCT TTCTTGGTGT AACTATCTTT	2400
ATTCTAACTC TG	2412

(2) INFORMATION FOR SEQ ID NO: 63:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 7760 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 63:

CCGATTTGGT GGAATTTTTG TCTCATCATT TAGAAGGTGT TGCAAGAGCA GAGTTTACCT	60
TGGTGCTTCA TACCAAATTG GGAGAAGCCT CTGTTTGGC AAAATTTGTA GATGAAACAA	120
AGGATGAATG GATTTAGGA ACAGTTGCTG GTGCCAATAC CTTATTGGTT ATTTGTCGAG	180
ATCAGCACGT TGCCAAACTC ATGGAAGATC GTTGCTAGA TTTGATGAAA GATAAGTAAG	240
GTCTTGGAG TTGCTCTCAA GACTTATTT TGAAAAGGAG AGACAGAAAA TGGCGATAGA	300
AAAGTTATCA CCCGGCATGC AACAGTATGT GGATATTAAG AAGCAATATC CAGATGCTTT	360

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TTTGCTCTT CGGATGGGTG ATTTTTATGA ATTATTTTAT GAGGATGCCG TCAATGCTGC	420
GCAGATTCTG GAAATTCCT TAACGAGTCG CAACAAGAAT GCCGACAATC CGATCCCTAT	480
GGCGGGTGT CCCTATCATT CTGCCCAACA GTATATCGAT GTCTTGATG AGCAGGGTTA	540
TAAGGTGGCT ATCGCAGAGC AGATGGAAGA TCCTAAACAA GCAGTTGGGG TTGTTAACAC	600
AGAGGTGTT CAGGTCTTA CGCCAGGGAC AGTGGTCGAT AGCAGTAAGC CGGACAGTCA	660
GAATAATTTC TTGGTTCCA TAGACCGCGA AGGCAATCAA TTTGGCCTAG CTTATATGGA	720
TTTGGTACG GGTGACTTTT ATGTGACAGG TCTTTGGAT TTCACGCTGG TTTGTGGGA	780
AATCCGTAAC CTCAAGGCTC GAGAAGTGGT GTGGGTTAT GACTTGCTG AGGAAGAAGA	840
ACAAATCCTC AGCCGCCAGA TGAATCTGGT ACTCTCTTAT GAAAAAGAAA GCTTTGAAGA	900
CCTTCATTAA TTGGATTTCG GATTGGCAAC GGTGGACCAA ACGGCATCTA GTAAGCTGCT	960
CCAGTATGTT CATCGGACTC AGATGAGGGG ATTGAACCAC CTCAAACCTG TTATCCGCTA	1020
CGAAATTAAG GATTCTTGC AGATGGATTA TCGGACCAAG GCTAGTCTGG ATTTGGTTGA	1080
GAATGCTCGC TCAGGTAAGA AACAAAGGCAG TCTTTCTGG CTTTTGGATG AAACCAAAAC	1140
GGCTATGGGG ATGCGTCTCT TGCGTTCTTG GATTCATCGC CCCTTGATTG ATAAGGAACG	1200
AATCGTCCAA CGTCAGAAAG TAGTGCAGGT CTTTCTCGAC CATTCTTTG AGCGTAGTGA	1260
CTTGACAGAC AGTCTCAAGG GTGTTTATGA CATTGAGCGC TTGGCTAGTC GTGTTTCTTT	1320
TGGAAAAACC AATCCAAGG ATCTCTTGCA GTTGGCGACT ACCTTGTCTA GTGTGCCACG	1380
GATTGCGTGC ATTMTAGAAG GGATGGAGCA ACCTACTCTA GCCTATCTA TCGCACAACT	1440
GGATGCAATC CCTGAGTTGG AGAGTTTGAT TAGCGCAGCG ATTGCTCCTG AAGCTCCTCA	1500
TGTGATTACA GATGGGGAA TTATCCGGAC TGGATTTGAT GAGACTTTAG ACAAGTATCG	1560
TTGCGTCTC AGAGAAGGG A CTAGCTGGAT TGCTGAGATT GAGGCTAAGG AGCGAGAAA	1620
CTCTGGTATC AGCACGCTCA AGATGACTA CAATAAAAG GATGGCTACT ATTTTCATGT	1680
GACCAATTG CAACTAGGAA ATGTGCCAGC TCACTTTTC CGCAAGGCAG CGCTGAAAAA	1740
CTCAGAACGC TTTGGAACCG AAGAATTAGC CCGTATCGAG GGAGATATGC TTGAGGCCCG	1800
TGAGAAGTCA GCCAACCTCG AATACGAAAT ATTATGCGC ATTCTGTGAAG AGGTCGGCAA	1860
GTACATCCAG CGTTTACAAG CTCTAGCCCA AGGAATTGCG ACGGTTGATG TCTTACAGAG	1920
TCTGGCGGTT GTGGCTGAAA CCCACCATTT GATTGACCT GAGTTGGTG ACGATTCAACA	1980
AATTGATATC CGGAAAGGGC GCCATGCTGT CGTTGAAAG GTTATGGGG CTCAGACCTA	2040
TATTCCAAAT ACGATTCAAGA TGGCAGAAGA TACCAAGTATT CAACTGGTTA CAGGGCCAAA	2100

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CATGAGTGGG AAGTCTACCT ATATGCGTCA GTTAGCCATG ACGGCGGTTA TGGCCCAGCT	2160	
GGGTTCTAT GTTCCTGCTG AAAGCGCCCA TTTACCGATT TTTGATGCGA TTTTTACCG	2220	
TATCGGAGCA GCAGATGACT TGTTTCCGG TCAGTCAACC TTTATGGTGG AGATGATGGA	2280	
GGCCAATAAT GCCATTCGC ATGCGACCAA GAACTCTCTC ATTCTCTTG ATGAATTGGG	2340	
ACGTGGAACT GCAACTTATG ACGGGATGGC TCTTGCTCAG TCCATCATCG AATATATCCA	2400	
TGAGCACATC GGAGCTAAGA CCCTCTTTGC GACCCACTAC CATGAGTTGA CTAGTCTGGA	2460	
GTCTAGTTTA CAACACTTGG TCAATGTCCA CGTGGCAACT TTGGACCAAG ATGGGCAGGT	2520	
CACCTTCCTT CACAAGATTG AACCGGGACC AGCTGATAAA TCTACGGTAT CCATGTTGCC	2580	
AAGATTGCTG GCTTGCAGC AGACCTTTTA GCAAGGGCGG ATAAGATTTT GACTCAGCTA	2640	
GAGAACAGAAG AACAGAGAG TCCTCCTCCC ATGAGACAAA CTAGTCTGT CACTGAACAG	2700	
ATTTCACTCT TTGATAGGGC AGAAGAGCAT CCTATCCTAG CAGAATTAGC TAAACTGGAT	2760	
GTGTATAATA TGACACCTAT GCAGGTTATG AATGCTTAG TAGAGTTAAA ACAGAAACTA	2820	
TAAACCAAG ACTCACTAGT TAATCTAGCT GTATCAAGGA GACTTCTTTG ACAATTCTCC	2880	
ACTTTTTGTC TAGATAACA TCACACAAAC AGAATGAAAA GGAGCTGACG CATTGTCGCT	2940	
CCCTTTGTC TATTTTTAA GGAGAAAGTA TGCTGATTCA GAAAATAAAA ACCTACAAAGT	3000	
GGCAGGCCCT GGCTTCGCTC CTGATGACAG GCTTGATGGT TGCTAGTTCA CTTCTGCAAC	3060	
CGCGTTATCT GCAGGAAGTC TTAGGCGCCC TCCTTACTGG GAAATATGAA GCTATTATA	3120	
GTATCGGGGC TTGGTTGATT GGTGTGGCCG TAGTCGGTCT AGTTGCTGGT GGACTCAATG	3180	
TTGTCCCTCGC AGCCTATATT GCCCAAGGAG TTTCATCCGA CCTTCGGGAG GATGCCCTCC	3240	
GTAAAATTCA AACCTTTCT TATGCTGATA TTGAACAATT TAATGCGGGA AATCTAGTCG	3300	
TTCGAATGAC AAATGATATC AACCAGATTC AGAACGTTGT CATGATGACC TTCCAAATT	3360	
TTTCAGACT TCCCCCTTG TTCATCGTT CGTTTATCCT AGCGGTTCAA ACCTTACCTT	3420	
CTCTGTGGTG GGTGATTGTT CTCATGGTAG TCTTGATTTT TGTTTGACT GCTGTCATGA	3480	
TGGGAATGAT GGGGCCTCGT TTTGCCAAGT TTCAAACCTT TCTTGAGCGC ATCAATGCCA	3540	
TTGCCAAGGA AAATTACGT GGCGTTCGTG TGGTCAAGTC CTGGTCCAA GAAAAGAGC	3600	
AATTTGCTAA GTTTACAGAG GTCTCAGACG AGCTTCTTGG TCAAAACCTT TACATTGGTT	3660	
ATGCCCTTTTC AGTAGTGGAA CCCTTTATGA TGTTGGTTGG TTACGGGGCG GTCTTCCCT	3720	
CTATTGCGCT GGTCGCGGGA ATGGTTCACT CGGATCCGTC TGTTGTTGGT TCCATCGCTT	3780	
CTTTTGTAA TTACCTAAGC CAGATTATCT TTACCATTTGT TATGGTTGGA TTTTTGGGAA	3840	
ATTCCTGTCAG CGGTGCCATG ATTCCATGC GTCGTATTG AGAAATTCTT GACGCAGAGC	3900	

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CAGCTATGAC CTTCAAGGAT ATCCCAGATG AAGAGTTGGT TGGAAAGTCTT AGCTTTGAAA	3960
ATGTGACCTT TACCTATCCA ATGGACAAGG AACCGATGCT GAAAGATGTG AGCTTTACTA	4020
TTGAACCTGG TCAAATGGTT GGTGTAGTTG GAGCGACTGG TGCAAGGAAAG TCAACCTTGG	4080
CTCAATTGAT TCCACAGTC TTTGATCCAC AGGACGGGGC CATTAAAATC GGTGGCAAGG	4140
ATATTGAGA AGTGAGTGA GGAACCCCTGC GTAAAACAGT TTCCATCGTT CTCCAACGTG	4200
CCATTCTTT TAGTGGAACG ATTGCAGATA ACTTGAGACA GGGGAAGGGG AATGCTACTC	4260
TATTTGAAAT GGAGCGCGCA GCCAATATTG CCCAGGCTAG TGAATTCAATT CATCGTATGG	4320
AGAAAACCTT TGAAAGTCCA GTTGAAGAAC GGGGAACCAA TTTCCTCTGGT GGACAAAAAC	4380
AAAGGATGTC GATTGCGCGT GGGATTGTCA GCAATCCACG TATTCTGATT TTGATGATT	4440
CGACCTCAGC CTTGGATGCC AAATCAGAGC GCTTGGTGCA AGAAGCTTG AATAAGGACT	4500
TGAAGGGAC GACAACCATT ATTATTGCTC AAAAATTAG CTCGGTTGTC CATGCAGACA	4560
AGATCTTGGT TCTAAATCAA GGACGATTGA TTGGTCAAGG TACGCATGCA GACTTGGTTG	4620
CCAACAATGC CGTTTACCGT GAAATCTATG AAACACAGAA ATGAAAGACA AACTATAAGA	4680
AAAGTCAATA GTTTTATCTA AACTATTTCT TATTTCAATT TGATGATTG GCGATGATT	4740
TAGAGCACGG CAAAAAGCCC TTGAAAAAGT CCATTTTTTC AAAGGTAATC CTGTGTTAAT	4800
TTCAGAAATT ACATCACTTT TTGTTCGTCA AATGGCAGCT CTTTTTTTAG GATATAAAAC	4860
AGGGTTCGGA TAACTTTTT TGCAAGGTGG ATGATGGCTA CATTGTAATG TTTTCCTTGT	4920
TCTAATTAG TCTTAAGATA GGCCTTAAAAA GCAGGGAAA AGCGAGGGCA TGCTTTGGCA	4980
GCTTGTATGA GTACCTACCG CAGATGAGGG GAACTCCGTT TGACCATTTCT TCCTGCTAAA	5040
TCAATCTGAT CTGACTGATA AATAGAAGAA TCCAGTCCAG CGAAAGCTTG TAATTGAGCA	5100
GGATTATCAA AGGCATGAAT ATTCGAATC TCAGCTAAA TGACCGCCCC TAAACGATCC	5160
CCAATCCCAG TAACCGTCGT GATGACCGAG TTGAACCTCAG CCATCAAGTC ATTGACACAT	5220
GTTCCTGCCT TGTCAATGAG CCTCTTGAA TGTTTGATGT TTTCATTACA CGAGATAAAA	5280
CGTCTATGCG TTATCAAAC TATTACCAAT TAAAACAAAA AGCTGTTGGT AGATCCTTTC	5340
GGAAATTGTC AAGCGATTGG AGGAAATGAA CTAATCCACA GCGGCTTATT CCAAGTATAC	5400
CACTTGGGCT TTGGCAGTAG CTAACTGCGC TAAATATAAT ATAAGGAGGA GTAAAATGAA	5460
GACAGTCAA TTTTTTGGC ATTATTTAA GGCTACAAG TTCTCATTTG TAGTTGTCAT	5520
CCTGATGATT GTTCTGGCGA CTTTTGCCCA AGCCCTCTTT CCAGTCTTT CTGGACAAGC	5580
GGTGACGCAG CTAGCCAATT TAGTTCAAGC TTATCAAAT GGCAATCCAG AACTTGTATG	5640

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GCAAAGCCTA	TCAGGAATCA	TGGTCAATCT	TGGCCTGCTG	TTTTGGTTC	TATTTATCTC	5700
TAGTGTAATA	TACATGTGTC	TCATGACGCG	CCTGATTGCA	GAATCGACCA	ACGAGATGCG	5760
CAAAGGCCCTC	TTTGGTAAGC	TTGCTCAGTT	GACGGTTCT	TTCTTGACC	GTCGACAAGA	5820
TGGCGATATC	CTGTCCTATT	TTACCCAGTGA	TTTGGATAAT	ATCCTCCAAG	CCTTTAACGA	5880
AAGCTTGATT	CAGGTCATGA	GCAATATTGT	TTTATACATT	GGTCTGATTC	TTGTCATGTT	5940
TTCGAGAAAT	GTGACGCTGG	CTCTCATCAC	CATTGCCAGC	ACCCCATTGG	CTTTCCCTTAT	6000
GCTGATTTTC	ATCGTGAAA	TGGCACGCAA	ATACACCAAC	CTCCAGCAGA	AAGAGGTAGG	6060
GAAGCTCAAC	GCCTATATGG	ATGAGAGCAT	CTCAGGCCAA	AAAGCCGTGA	TTGTGCAAGG	6120
AATTCAAGAG	GATATGATGG	CAGGATTTCT	TGAACAAAAT	GAGCGCGTGC	GCAAGGCAAC	6180
CTTTAAAGGA	AGAATGTTCT	CAGGAATTCT	TTTCCCTGTC	ATGAATGGGA	TGAGCCTGAT	6240
TAATACAGCC	ATCGTCATCT	TTGCTGGTTC	GGCTGTACTT	TTGAATGATA	AGTCTATTGA	6300
AACAAGTACA	GCCCTAGGTT	TGATTGTTAT	TTTGCACAA	TTTCACAGC	AGTACTACCA	6360
GCCTATTATC	CAAGTTGCAG	CGAGTTGGGG	AAGCCTTCAG	TTGGCCTTTA	CTGGAGCTGA	6420
ACGAATTCA	GAAATGTTG	ATGCAGAGGA	GGAAATCCGA	CCTGAAAAGG	CTCCAACCTT	6480
CACTAAGTTG	CAAGAAAGTG	TTGAAATCA	TCATATCGTT	TTTCATACT	TGCGCTGATAA	6540
ACCTATTTG	AAAGATGTCA	GCATTTCTGC	CCCTAAAGGC	CAGATGACAG	CAGTTGTTGG	6600
GCCGACAGGT	TCAGGAAAAA	CGACTATTAT	GAACCTCATC	AATCGCTTTT	ATGATGTTCA	6660
TGCTGGTGGT	ATTTATTTG	ATGGTAAAGA	CATTGCGGTC	TATGACTTAG	ATAGTCTTAG	6720
AAGCAAGGTG	GGAATTGTAT	TGCAAGATTG	GGTCTTGT	AGCGGAACGA	TTAGAGACAA	6780
TATCCGATTT	GGTGTGCGAG	ATGCTAGTCA	GGAAATGGTT	GAGGTAGCAG	CAAAGCAAC	6840
CCACATTCAC	GACTATATCG	AAAGTTGCC	TGATAAGTAC	GATACTCTTA	TTGATGATGA	6900
CCAGAGCATC	TTTCAACAG	GGCAGAAGCA	ATTGATTCA	ATCGCTCGAA	CCCTGATGAC	6960
AGATCCAGAA	GTTCTCATTC	TCGATGAAGC	AACTCAAAC	GTAGATACGG	TGACAGAAAG	7020
CAAGATTCA	CATGCCATGG	AGGTGGTTGT	ACGAGGTAGA	ACTAGTTCG	TCATTGCCCA	7080
CCGCTTGAAA	ACCATTCTCA	ATGCAGATCA	GATTATTGTC	CTTAAAGATG	GAGAAGTCAT	7140
TGAACCGTGGT	ACCACCATG	AACTTTGAA	GCTAGGTGGC	TTTTATTCA	AACTCTATCA	7200
CAATCAATT	TTTTCGAAT	AAGAAAGAAG	TTGTCCTATG	TGGGCAGCTT	TTTCTTGTCC	7260
ATAAAAAAATG	TTTATCACAG	CCTAAAAAAA	AAACATATTAG	ACGAAAGTCA	TTTGAGTGA	7320
TATGATAGGA	CTATCGTTAG	CATTGAAAG	GAGAGGCATC	ATGGCTAGAA	CGGTTGTAGG	7380
AGTTGCTGCA	AATCTATGTC	CCGTAGACGC	AGAAGGCAA	ATCATTCA	CATCTGTATC	7440

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TTGTAGATTC GCAGAGATCA TTCGTCAAGT CGGTGGTCTC CCTTTAGTCA TTCCCTGTTGG	7500
TGATGAGTCA GTTGTACGTG ATTATGTGGA AATGATTGAC AAACTCATTT TGACAGGAGG	7560
CCAAAATGTT CATCCTCAGT TTTATGGAGA GAAAAAGACC GTCGAGAGCG ATGATTACAA	7620
TCTGGTCCGT GACGAATTTG AATTGGCACT CTTGAAGGAA GCGCTTCGTC AGAATAAACCC	7680
AATTATGGCA ATCTGTCGCG GTGTCCAATCT TGTCATGTT GCCTTTGGTG GAACCCCTCAA	7740
TCAAGAAATC GAAGGTCAGG	7760

(2) INFORMATION FOR SEQ ID NO: 64:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 2723 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 64:

GAGGTTTTAA TTCACATTAC TCTsCCGTAT CTTTATTTAA AATGAATTCT TTTACGGTTG	60
TATTTCTTGC AAAATCTTTT ACAACAATCT TAATGTTTAG TGTCTTGCT ATTATTTGTT	120
TAATATCATT AAATGATGTA TATTCTTTTC CATTATATA AATATGTTGT TCTTGAATCT	180
CACCATCGAA TCCATTATTT CTTTTATCAT TGATGTTAAA GACTACAGAT TTTCCATCAG	240
CATATTCGAT ACTAGTATTT CCCTTAGGAT CAATGTTAAC TTGCGGGTTA ACATTATCAT	300
ATAAAAACTG ATAGTGGACT CCAACTGCTT TAGCATTCAA ATCGCTATAG CCAGTTGAA	360
GATAAACATT TCCATCCATA TCTGTTACCT TATCTGGAAA TCCGTTTGCT TTATAGTCTT	420
TCATTCCCCA GTCCATGATG TCACCGTCTT TAACATTCAAG CTTAATATTA AAATCTCTAG	480
TGTTATCAAT GTGTAATCT CCGTAGATTA AATAATTATC TACAACCGAT TCATTAACCTC	540
TCAATTCCCA CTTAAAACCA CCCTTATCAG AAATCTTACC TCTTAAATAA AATTCTGGAT	600
TTCGTACATA AATTCTTATTA GATTAGATG GATTAAGTA GTTCTTATCC ATTGAAAGGT	660
TTACTGGTTT GGTATCAATA AATAACATGG AGCCATCTTC TTTTATAGCT TCTACATTGA	720
ACTTATCCTC TCCAGTGTAT TCTTTATCAT CCTTACAAA TAATACAAAGT TTAGAAGAAAT	780
CTGTCACAAG ATTTCCGTCT TTATCGATAG CTTCCCTTT ATCGTTCATT TTAAATGAA	840
ACACTTGATA CCTTATAATG TTAAAGCCGT CCAAAGCCGA CATTAAATACA GATTGGGTAC	900
TTCTTCCATC TTCAACATTT CTACTATCAG CATAAATTGT TGTTCTGAA AGGGCTCTTA	960
GATTAGGATT GGCCCTTTGT ATTCTTGCTA TATCTCCCTT GCTATAGACT CCATTTCCCTT	1020

538

CTAACATATC CGTTTTCCA GGATTATAGG TAGTCAC TTT TAGTCATAG CCTTTCTTA	1080
GAATGATATT ATCCTTAAAC AGATATTGTT GTTTTCTGA ATCAGAATAG ATTTTACCAAG	1140
ATTCCATTGT AGTTAAATTG TCTGGTTGT TTTTGAAAG ATTCCTTCC CCTAATTCTA	1200
TGACATTCCC ATAACTTGAT ACATAGGGAT ATTCTGATT AGTTTCTTA ATTTTTTCAG	1260
GCATTCTAAAT TTAAATTCA GCTTTTCT GATCATTATC TTAAACAAAT AATCTCATAT	1320
CTCCTGAAA AGCTAATCCA TCCACAATAT CATTAAATT AGCGTATAGA TCAAATGTCA	1380
TCGTTTTGA GTGAAATCA TACTGGTCG CTTGATTTC TATAGATTAA TAGTTATTCC	1440
CATAATATAC CTTGGCATT TTAGAAACAT TACTTATCTT TCCAAGAATT TCAAAGTGTG	1500
CATCTTCTAGA CGGACTTAGA ACCCATAAA TTTTGATTGATTT GATTCGTCA AGTTTCTCAG	1560
TTTCATATTC TAGATCAGTC CCATCATCGT AGGCTATTAT ATTCCTTTA TCATCGTATT	1620
TATAATCGTA TTCCCTCATT CTCTTACCAAG TTTCACTTGT AAAATCATCA ACTTCTCTAA	1680
ATTTCTTTT AATGAGTTTC TTAAAGTCTT TATTTCAAA GTCTCTAATT GTGAAATAT	1740
TTCTATCAAT AGTAAAACTA GATTTTCTT TAATAGACTC TTCATTTCT TGATGATGAT	1800
GTTCTACCCC AGTTGTATCT TTTTTAGAC TACCTCTTT TCCATTCTT AAATTTTAA	1860
ATTTAGATTC TGCAATCTCG CCAAGCTTT GATATTAGA TGAATCTTGA TCAGGATCTA	1920
CTAGATAATA GGAAATCATC CCCTTTCTAT CAGCCTGATT AGCAAATTAA ATTCTATGAA	1980
TCTTTGTGAA ATTGCTAGAA CCATCTAATG CAATGACTTC AATGATTTT CCCCTTAAAT	2040
CTCCCCGACC TTAAATTCA TAAATGGTAT TTCCGTCTT ATCAAGTTTT CTATTCTTCA	2100
CTTGACCCCTC ACCTGCGTAA GTTACTTCAA GATTTTTTC AACCTCTCCA TCTTCATTAA	2160
CAAGAGCGGC GCCAGCATAAC CAAACTTCGT TCGCAATCTC GTCAAATTAA TCAGGATGTT	2220
CTTTTGATC TCTCGAAAT AGCGTTCTAT TCTTATACTG ATCTTTTACCC TTATGATAAG	2280
TATCCTTGT AATCAACTTA ATTTTTCAAG GATTTGAAAA ATCAACCGAA ACAATCTTAG	2340
GGGCGGTGTT ATCAATTAAACAGGAATAT AGGAAACCTG CCATGGGTAA TCTTAGTTA	2400
ATCTATATTT AAATTATAG AAATATTGAC CTTCCGCAAT CGGTTCAAAT TGACCTCTTA	2460
TCTTAGTAGC AGGATCTTGA TTATCCTTAC TTTCTGGTGC ATTTCTTCT CTACCTCTAG	2520
GATTATAGAT GAGTCCATCC CACTTCAGT CACCCCAAC TTTTAGTTA GATGATTGAA	2580
TTCCCTTGC ATCATTGCTT TTAGAATTAA AAATTCTCTT AATAAAGTGT TCTCTCGAAA	2640
TGACTTTAA GTCTCTTGA TTTCTCCCT CTTTATTGTT ATTTACTATT GAAATCAATC	2700
CTTCTCTGC ACTTCTTAAT ACA	2723

(2) INFORMATION FOR SEQ ID NO: 65:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 11831 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 65:

AAAAAAAGTGG	GAATGACTCA	AATCTTCACT	GAAGCTGGCG	AATTGATCCC	TGTAACAGTT	60
ATTGAAGCAA	CTCCAAACGT	TGTTCTTCAA	GTTAAAACGT	TTGAAACAGA	CGGATAACAC	120
GCTATCCAAG	TTCGGTTTCGA	TGACAAACCC	GAAGTATTGA	GCAACAAACC	TGCTAAAGGA	180
CATGTAGCGA	AAGCTAACAC	GGCTCCTAAG	CGCTTCATTG	GTGAATTCAA	AAACGTTGAA	240
GGCTTGAAG	TTGGTGCTGA	AATTACAGTT	GAAACATTG	CAGCTGGAGA	CGTTGTTGAC	300
GTAACGGTA	CTTCTAAAGG	TAAAGGTTTC	CAAGGTGTTA	TCAAACGCCA	CGGACAATCA	360
CGTGGACCAA	TGGCTCACGG	TTCTCGTTAC	CACCGTCGTC	CAGGTTCTAT	GGGGCCTGTT	420
GCACCTAAC	CGCTATTCAA	AGGTTAAAAC	CTTGCAGGAC	GTATGGGTGG	CGACCGCGTA	480
ACAATTCAA	ACCTTGAACT	TGTACAAGTT	GTTCCAGAAA	AGAACGTTAT	CCTTATCAA	540
GGTAACGTAC	CAGGTGCTAA	GAAATCTTT	ATCACTATCA	AATCAGCAGT	AAAGCTGGT	600
AAATAATAAA	GAAAGGGAA	ATCAGTCACA	ATGCCAACG	TAACATTATT	TGACCAAAC	660
GGTAAAGAAG	CAGGTGCTAA	GAAATCTTT	ATCACTATCA	AATCAGCAGT	AAAGCTGGT	720
TCAGTTGTGT	TTGATGTAAT	CATCAGCCAA	CGCGCAAGCC	TTCGTCAAGG	AAACACACGCT	780
GTTAAAACCC	GCTCTGCAGT	ATCAGGTGGT	GGACGCCAAC	CATGGCGTCA	AAAAGGAAC	840
GGACGTGCTC	GTCAGGTTTC	TATCCGCTCA	CCACAATGGC	GTGGTGGTGG	TGTTGTCTTC	900
GGACCAAAC	CACGTTCATCA	CGGCTACAAA	CTTCCACAAA	AAGTCGTCG	CCTAGCTCTT	960
AAATCAGTTT	ACTCTGAAA	AGTTGCTGAA	AACAAATTG	TAGCTGTAGA	CGCTCTTCA	1020
TTTACAGCTC	CAAAACTG	TGAATTGCA	AAAGTTCTTG	CAGCATTGAG	CATCGATTCT	1080
AAAGTTCTTG	TTATCCTTGA	AGAAGGAAAT	GAATTGCGAG	CTCTTTGAGC	TGCTAACCTT	1140
CCAAACGTGA	AAAGTGCAAC	TGCTACAAC	GCAAGTGTTC	TTGACATCGC	AAATAGCGAC	1200
AAACTTCTTG	TCACACAAGC	AGCTATCTCT	AAAATCGAGG	AGGTTCTTGC	ATAATGAATT	1260
TGTATGATGT	TATCAAAAAA	CCTGTCATCA	CTGAAAGCTC	AATGGCTCAA	CTTGAAGCAG	1320
GAAAATATGT	ATTTGAAGTT	GACACTCGTG	CACACAAACT	TTTGATCAAG	CAAGCTGTTG	1380
AAGCTGCTTT	CGAAGGTGTT	AAAGTTGCCA	ATGTTAACAC	AATCAACGTA	AAACCAAAAG	1440

540	
CTAAACGTGT TGGACGTTAC ACTGGTTTA CTAACAAAAC TAAAAAAGCT ATCATCACAC	1500
TTACAGCTGA TTCTAAAGCA ATCGAGTTGT TTGCTGCTGA AGCTGAATAA TCTAAGGAGG	1560
AAATATCGTG GGAATTCTGTG TTTATAAACC ACAACAAAAC GGTCGCCGTA ATATGACTTC	1620
TTTGGATTTG GCTGAAATCA CAACAAGCAC TCCTGAAAAA TCATTGCTTG TTGCATTGAA	1680
GAGCAAGGCT GGTCGTAACA ACAACGGTCG TATCACAGTT CGTCACCAAG GTGGTGGACA	1740
CAAACGTTTC TACCCTTTGG TTGACTCAA ACCTAATAAA GACAACGTTG AAGCAGTTG	1800
TAAAACAATC GAGTACGATC CAAACCGTTC TGCAAACATC GCTCTTGTAC ACTACACTGA	1860
CGGTGTGAAA GCATACATCA TCGCTCCAAA AGGTCTTGAA GTAGGTCAAC GTATCGTTTC	1920
AGGTCCAGAA GCAGATATCA AAGTCGAAA CGCTCTTCCA CTTGCTAACAA TCCCAGTTG	1980
TACTTTGATT CACAACATCG AGTTGAAACC AGGTCTGTTG GTTGAATTGG TACGTGCTGC	2040
TGGTGCATCT GCTCAAGTAT TGGGTTCTGA AGGTAAATAT GTTCTTGTTC GTCTTCAATC	2100
AGGTGAAGTT CGTATGATTC TTGGAACCTTG CCGTGCTACA GTTGGTGTG TCGGAAACGA	2160
ACAACATGGA CTTGTAAACC TTGGTAAAGC AGGACGTAGC CGTTGGAAAG GTATCCGCC	2220
AACAGTTCGT GGTTCTGTA TGAACCCCTAA CGATCACCCA CACGGTGGTG GTGAAGGTA	2280
AGCACCAGTT GGTCGTAAG CACCATCTAC TCCATGGGGC AAACCTGCTC TTGGTCTTAA	2340
AACTCGTAAC AAGAAAGCGA AATCTGACAA ACTTATCGTT CGTCGTCGCA ACGAGAAATA	2400
ATATTAACACT AGTCGCTTA GCAACTAGTA AATCCGCCAG CTCGGTAGCG CTCCATAGGA	2460
GTGCAAGCCG CTGTGGTACA ACATTTAAAG GAGAAAATAT AAAATGGGA CGCAGTCTTA	2520
AAAAAGGACC TTTCGTCGAT GAGCATTGTA TGAAAAAAGT TGAAGCTCAA CCTAACGACG	2580
AAAAGAAAAA AGTTATTAAA ACTTGGTCAC GTCGTTCAAC GATCTTCCCA AGTTTCATTG	2640
GTTACACTAT TGCAGTTTAT GACCGACGTA AACACGTACC TGTTTACATC CAAGAAGACA	2700
TGGTAGGCCA CAAACTTGGT GAATTGAC CAACTCGTAC TTACAAAGGT CACGCTGCAG	2760
ACGACAAGAA AACACGTAGA AAATAAGGAG AACATAAATG GCAGAAAATTA CTTCAGCTAA	2820
AGCAATGGCT CGTACAGTAC GTGTTTCACC TCGTAAATCA CGTCTTGTTC TTGATAACAT	2880
CCGTGGTAA AGCGTAGCCG ATGCAATCGC AATCTTGACA TTCACTCCAA ACAAAGCTGC	2940
TGAAATCATC TTGAAAGTTT TGAACTCAGC TGTAGCTAAC GCTGAAAACA ACTTTGGTTT	3000
GGATAAAAGCT AACCTGGTAG TATCTGAAGC ATTCTGCAAAC GAAGGACCAA CTATGAAACG	3060
TTTCCGTCCA CGTGCAGAAAG GTTCAGCTTC ACCAATCAAC AACACGTACAG CTCACATCAC	3120
TGTAGCTGTT GCAGAAAAT AAGGAGGTAA AATCGTGGGT CAAAAAGTAC ATCCAATTGG	3180
TATCGTGTC GGCATCATCC GTGATTGGGA TGCCAAATGG TATGCTGAAA AAGAATACGC	3240

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GGATTACCTT CATGAAGATC TTGCAATCCG TAAATTGTT CAAAAAGAAC TTGCTGACGC	3300
AGCAGTTTC ACTATTGAAA TCGAACGCGC AGTAAACAAA GTTAACGTTT CACTTCACAC	3360
TGCTAAACCA GGTATGGTTA TCGGTAAAGG TGTTGCTAAC GTTGATGCaC TCCGTGCAA	3420
ACTTAAACAAA TTGACTGGAA AACAAAGTACA CATCAACATC ATCGAAATCA AACAAACCTGA	3480
TTTGGATGCT CACCTGTAG GTGAAGGAAT TGCTCGTCAA TTGGAGCAAC GTGTTGTTT	3540
CCGTCGTGCA CAAAAACAAG CAATCCAACG TGCAATGCGT GCTGGAGCTA AAGGAATCAA	3600
AACTCAAGTA TCAGGTCGTT TGAACGGTGC AGATATCGCC CGTGTGAAG GATACTCTGA	3660
AGGAACATGTT CCGCTTCACA CACTTCGTGC AGATATCGAT TACGCCGGG AAGAACAGA	3720
TACTACATAC GGTAAACCTG GTGTTAAAGT ATGGATCTAC CGTGGTGAAG TTCTTCCAGC	3780
TCGTAAAAAC ACTAAAGGAG GTAAATAACC AATGTTAGTA CCTAAACGTG TTAAACACCG	3840
TCGTGAGTTC CGTGGAAAAA TGCGCGGTGA AGCAAAAGGT GGAAAAGAAC TAGCATTCCG	3900
TGAATACGGT CTTCAAGCTA CAACTAGCCA CTGGATCACT AACCGCCAAA TCGAAGCTGC	3960
TCGTATCGCC ATGACTCGTT ACATGAAACG TGGTGGTAA GTTTGGATTA AAATCTTCCC	4020
ACACAAATCA TACACTGCTA AAGCTATCGG TGTGCGTATG GGATCTGGTA AAGGGGCACC	4080
TGAAGGTTGG GTAGCACAG TTAAACGTGG TAAAGTGTATG TTCGAAATCG CTGGTGTATC	4140
TGAAGAGATT GCACGTGAAG CGCTTCGACT TGCTAGCCAC AAATTGCCAG TTAAATGTAA	4200
ATTCGTAAAA CGTGAAGCAG AATAAGGAGA AGGCATGAAA CTTAATGAAG TAAAAGAATT	4260
TGTTAAAGAA CTTCGTGGTC TTTCTCAAGA AGAACTCGCG AAGCGCGAAA ACGAATTGAA	4320
AAAAGAATTG TTTGAACCTTC GTTCCAAGC TGCTACTGGT CAATTGGAAC AAACAGCTCG	4380
CTTGAAAGAA CTTAAAAAAC AAATCGCTCG CATCAAAACA GTTCAATCTG AAGCGAAATA	4440
ATAGACTAGG GAAGGAGAAA TTCAATGGA ACGCAATAAT CGTAAAGTTC TTGTTGGACG	4500
TGTTGTATCT GACAAAATGG ACAAGACAAT CACAGTTGTA GTTGAACAA AACGTAACCA	4560
CCCAGTCTAT GGTAAACGTA TTAACTACTC TAAAAATAC AAAGCTCATG ATGAAAACAA	4620
TGTTGCCAAA GAAGGCCATA TCGTACGTAT CATGGAAACT CGCCCGCTTT CAGCTACAAA	4680
ACGTTTCCGT CTTGTAGAAG TTGTTGAAGA AGCGGTCATC ATCTAATCAA ACCTGAAAGG	4740
AGAAAACGTAA AATGATTCAA ACAGAAACTC GTTTGAAAGT CGCAGACAAC AGCGGTGCTC	4800
GCGAAATCTT GACTATCAA GTTCTTGGTG GTTCAGGACG TAAATTGCA AACATCGGT	4860
ATGTTATCGT GGCATCTGTA AAACAAGCTA CTCCCTGGTGG TGCGGTTAAA AAAGGTGACG	4920
TTGTTAAAGC AGTTATCGTT CGTACTAAAT CAGGTGCTCG TCGTGCTGAT GGTTCATACA	4980

542

TCAAATTGAGA CGAAAACGCCA GCAGTTATCA TCCGTGAAGA CAAAACTCCT CGCGAACAC	5040
GTATCTTGG CCCAGTTGCA CGTGAATTGC GTGAAGGTGG CTTCATGAAG ATCGTGTAC	5100
TTGCTCCAGA AGTACTTTAA TTTTTAGGAA CAAACTAGTC CCCTAGCTTC AAGCTAGGGT	5160
GCCCTTATGG GCGTAAGAAA AATCAAGGAG AACCTAATG TTTGTAAAAA AAGGCGACAA	5220
AGTTCCGCTA ATCGCTGGTA AAGATAAGGG AACAGAAGCT GTTGTCCCTTA CTGCCCTTC	5280
AAAAGTAAAC AAAGTTATCG TTGAAAGGTGT TAACATTGTT AAGAAACACC AACGTCCAAC	5340
TAACGAGCTT CCTCAAGGTG GTATCATCGA GAAAGAAGCA GCTATCCACG TATCAAACGT	5400
TCAAGTTTG GACAAAAATG GTGTAGCTGG TCGTGTGGA TACAAATTG TAGACGGTAA	5460
AAAAGTTCGC TACAACAAAA AATCAGGGCGA AGTGCTTGAT TAATCACGAA GGAAAGGAGA	5520
AGTATAATGG CAAATCGTT AAAAGAAAAA TATCTTAATG AAGTAGTTCC TGCTTTGACA	5580
GAACAATTCA ACTACTCATC AGTGATGGCT GTGCCCTAAAG TAGATAAGAT TGTTTGAAAC	5640
ATGGGTGTTG GTGAAGCTGT ATCAAACGCT AAAAGCCTTG AAAAGCTGC TGAAGAATTG	5700
GCACATTATCT CAGGTCAAAA ACCACTTATC ACTAAAGCTA AAAATCAAT CGCCGGCTTC	5760
CGTCTTCGTG AAGGTGTTGC GATCGGGTGC AAAGTTACCC TTCGTGGTGA ACGTATGTAC	5820
GAATTCTTGG ATAAATTGGT ATCAGTTCA CTTCCACGTG TACGTGACTT CCACGGTGTC	5880
CCAACAAAAT CATTGATGG ACGCGGGAAC TACACACTTG GTGTGAAAGA ACAATTAAATC	5940
TTCCCAGAAA TCAACTTCGA TGACGTTGAC AAAACTCGTG GTCTTGACAT CGTTATCGTA	6000
ACAAC TGCTA ACAC TGACGA AGAGTCACGT GCATTGCTTA CAGGCCTTGG AATGCCCTTT	6060
GCAAAATAAT ATAGGAGGTA AATCTAATGG CTAAAAAATC AATGGTAGCT AGAGAGGCTA	6120
AACGCCAAAA AATTGTTGAC CGTTATGCTG AAAAACGTGC TGCATTAAAG GCGGCAGGGG	6180
ACTACGAAGG TTTATCTAAA TTACCTCGCA ACGCCTCACC GACTCGTTA CATAATCGTT	6240
GTAGGGTTAC GGGGCCCA CATTCAAGTTT ACCGCAAATT TGGTCTGAGT CGTATCGCTT	6300
TTCGCCAACT TGCGCATAAA GGTCAAATTC CTGGTGTAAAC AAAAGCATCT TGGTAATTAA	6360
AGATATCAAG AGCGTCAAAA CTCCAAGTAA AAATAGGAA CTTGACGAAG AACTAAAGT	6420
TTCTAGGAAA GTTTATCTTT TTCACACAGA GTTTAGCCCG GGTTCAATTG GGCTTGCCAA	6480
TTTGAACACG AGCTACAGCT TTGGCAAAAA AGACCAATTG CTTTTGGAGC ATTGCTTCTG	6540
CATTAATTG TCTATTTTG CTCGTGCTGT TACGCTCTTT GTATCATGTA TTAACTAGCA	6600
AGTGCAACTT GCAAACACT AGTAAGAGGA GAAAAACAAA ATGGTTATGA CTGACCCAAT	6660
CGCAGACTTC CTAAC TCGTA TTCGTAATGC TAACCAAGCT AACACGAAG TACTTGAAAGT	6720
ACCTGCATCA AACATCAAAA AAGGGATTGC TGAAATCCTT AAACGCGAAG GTTTGTAAA	6780

AAACGTTGAA ATCATTGAAG ATGACAAACA AGGCAGTCATC CGTGTATTC TAAATACGG	6840
ACCAAAATGGT GAGAAAGTTA TCACTAACCTT GAAACGTGTT TCTAAACCAG GACTTCGTGT	6900
CTACAAAAAA CGTGAAGACC TTCCAAAAGT TCTTAACCGGA CTTGGAATTG CCATCCTTTC	6960
AACTTCTGAA GGTTTGCTTA CTGATAAAGA AGCACGCCAA AAGAATGTTG GTGGTGAGGT	7020
TATCGCTTAC GTTTGGTAAA ATCAAGATAC AAAGCTCGTA AAGAACAAAG CAAAATTAGG	7080
AAGTTGGAGA AGTTTGTGTTA CAAACAAGCC AACTTATCTA TTTTGACAG TTCTTAGAGC	7140
GTGTTCACTT CAGCTCTTGA ACTAAATAAG TATCTGAACC CCGTGAAAC TGCCCGTTCT	7200
GGCCTGACAA TTTAACAGGA GAAAATAAAC ATGTCACGTA TTGGTAATAA AGTTATCGT	7260
TTGCCTGCTG GTGTTGAACCT CGCTAACAT GACAACGTTG TAACGTAAA AGGATCTAAA	7320
GGAGAACCTTA CTCGTGAGTT CTCAAAAGAT ATTGAAATCC GTGTGGAAGG TACTGAAATA	7380
ACTCTTCACC GTCCAAACGA TTCAAAAGAA ATGAAAAC TCCACGGAAC TACTCGTGCC	7440
CTTTTGAAACA ACATGGTTGT TGGTGTATCA GAAGGATTCA AGAAAGAACT TGAAATCGGT	7500
GGGGTTGGTT ACCGTGCACA GCTTCAAGGA TCTAAACTTG TTTTGGCTGT TTGGTAAATCT	7560
CATCCAGACG AAGTTGAAGC TCCAGAAGGA ATTACTTTG AACTTCCAAA CCCAACAAACA	7620
ATCGTTGTTA GCGGAATTTC AAAAGAAGTA GTTGGTCAA CAGCTGCTTA CGTACGTAGC	7680
CTTGGTTCAC CAGAACCATATA TAAAGTAAA GGTATCCGTT ACGTTGGTGA ATTCTGTTCG	7740
CGTAAAGAAG GTAAAACAGG TAAATAATGT TGAGTGGTTG ATCATCAACC ACCAACCTAT	7800
TTTCCAACCTT TGTGCATAGC ACACGATTAA AACTAAAGA GGTGAAAAC GTGATTTCAA	7860
AACCAGATAAA AAACAAACTC CGCCAAAAC GCCACCGTCG CGTTCGCGGA AAACCTCTG	7920
GAACCTGCTGA TCGCCCACGT TTGAACGTAT TCCGTTCTAA TACAGGCATC TACGCTCAAG	7980
TGATTGATGA CGTAGCGGGT GTAACGCTCG CAAGTGCCTC AACTCTTGAT AAAGAAGTT	8040
CAAAAGGAAC TAAAATGAA CAAGCCGTTG CTGTCGGTAA ACTCGTTGCA GAACGTGCAA	8100
ACGCTAAAGG TATTCAGAA GTGGTGTTCG ACCGGCGGTGG ATATCTATAT CACGGACGTG	8160
TGAAAGCTTT GGCTGATGCA GCTCGTAAA ACGGATTGAA ATTCTAATAG GAGGACACTA	8220
GAAAATGGCA TTTAAAGACA ATGCAGTTGA ATTAGAAGAA CGCGTAGTTG CTGTCAACCG	8280
TGTTACAAAAA GTTGTAAAG GTGGACGTGCG TCTTCGTTTC GCAGCTCTTG TTGGTGTG	8340
TGACCCACAAT GGTCGCGTAG GATTTGGTAC TGGTAAAGCT CAAGAAGTTC CAGAACAAAT	8400
CCGTAAGCA GTAGATGATG CTAAGAAAAA CTTGATCGAA GTTCCTATGG TTGGAACAAAC	8460
AATCCACAC GAAGTCTTT CAGAACCGG TGAGCTAA GTATTGTTGA AACCTGCTGT	8520

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AGAAGGTTCT GGAGTTGCCG CTGGTGGTGC AGTCGTGCC	GTTGTGGAAT TGGCAGGTGT	8580
GGCAGATATT ACATCTAAAT CACTTGGTTC TAACACTCCA	ATCAACATTG TTCGTGCAAC	8640
TGTTGAAGGT TTGAAACAAT TGAAACGCCG	TGAAGAAATT GCTGCCCTTC GTGGTATTTC	8700
AGTTCTGAT TTGGCATAAG AAAGGGGATA	AAATGGCTCA AATTAAAATT ACTTTGACTA	8760
AGTCTCCAAT CGGACGCATT CCATCACAAC	GTAAAACGT TGAGCACTT GGACTTGGCA	8820
AATTGAACAG CTCTGTTATT AAAAGAAGATA	ACGCTGCTAT CCGTGGTATG ATCACAGCAG	8880
TATCTCACTT AGTAACAGTT GAAGAAGTAA	ACTAATGAaG TTTTAGGGGA TGTGCACTGT	8940
ACCATCCCCT AAAACTAGAT ATAGTCATCT	ATGATGACAT CGTATAGGCG AGTTGATGGG	9000
GGAGACAACC TTTCTCCCT TATCGCGCT	ACCATTTTAC AAAAGAGGAG AAAATAAAA	9060
TGAAACTTCA TGAATTGAAA CCTGCAGAAG	GTTCTCGTAA AGTACGTAAC CGCGTTGGTC	9120
GTGGTACTTC ATCAGGTAAC GGTAAAACAT	CTGGTGTGG TCAAAAGGT CAAAAAGCTC	9180
GTAGCGGTGG CGGAGGTCGC	CTTGGTTTG AAGGTGGACA AACTCCATTG TTCCGTCGTC	9240
TTCCAAAACG TGGATTCACT AACATCAACG	CTAAAGAATA CGCAATTGTG AACCTTGACC	9300
AATTGAACGT CTTTGAAGAT GGTGCTGAAG	TAACCTCCAGT TGTTCTTATC GAAGCAGGAA	9360
TTGTTAAAGC TGAAAAGTCA GGTATTAAAA	TTCTTGGTAA CGGTGAGTTG ACTAAGAAAT	9420
TGACTGTGAA AGCAGCTAA TTCTCTAAAT	CAGCTGAAGA AGCTATCACT GCTAAAGGTG	9480
GTTCAGTAGA AGTCATCTAA GAGAGGTGAC	CTATGTTTT TAAATTATTA AGAGAAGCTC	9540
TTAAAGTCAA GCAGGTTCGA	TCAAAAATT TATTTACAAT TTTTATCGTT TTGGTCTTT	9600
GTATCGAAC TAGCATTACA GTTCTGGTG	TGAATGCCAA TAGCTTGAAT GCTTTAAGTG	9660
GATTATCCTT CTTAACATG TTGAGCTTGG	TGTCGGGAA TGCCCTAAAA AACTTTCCGA	9720
TTTTGCCCC AGGAGTTAGT CCCTATATCA	CCGCTTCTAT TGTTGTCAA CTCTTGCAAA	9780
TGGATTTT ACCCAAGTT GTAGAGTGGG	GTAAACAAGG GGAAGTAGGT CGAAGAAAAT	9840
TGAATCAAGC TACTCGTTAT ATTGCTCTAG	TTCTCGCTTT TGTGCAATCT ATCGGGATTA	9900
CAGCTGGTT TAATACCTTG GCTGGAGCTC	AATTGATTAA AACTGTTTA ACTCCACAAAG	9960
TTTTCTGAC GATTGGTATC ATCTAACAG	CTGGTACTAT GATTGTCACT TGTTGGGTG	10020
AGCAAATTAC AGATAAGGGA TACGGAAACG	GTTTTCCAT GATTATCTT GCCGGGATTG	10080
TTCCCTCAAT TCCAGAGATG ATTCAAGGCA	TCTATGTGGA CTACTTTGTG AACGTCCCAA	10140
GTAGCCGTAT CACTTCATCT ATCATTTCG	TAATCATTGTT GATTATTACT GTATTGTTGA	10200
TTATTACTT TACAACCTTAT GTTCAACAAG	CAGAATACAA AATTCCAATC CAATATACTA	10260
AGGTTGCACA AGGTGCTCCA TCTAGCTCTT	ACCTTCCGTT AAAAGTAAAC CCTGCTGGAG	10320

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TTATCCCTGT TATCTTGCC AGTCGATTA CTGCAGCCTG CGGCTATTCT TCAGTTTTG	10380
AGTCCCACAG GTCATGATTG GGCTGGTA AGGGTAGCAC AAGAGATGTT GGCAACTACT	10440
TCTCCAATG GTATTGCCAT GTATCCTTG TTGATTATTC TCTTACATT CTTCTATAACG	10500
TTTGTACAGA TTAATCCTGA AAAAGCAGCA GAGAKCCTAC AAAAGAGTGG TGCCTATATC	10560
CATGGAGTTC GTCCGGTAA AGGTACAGAA GAATATATGT CTAAACTTCT TCGTCGTCTT	10620
GCAACTGTTG GTTCCCTCTT CCTTGGTGTG ATTTCCATT TACCGATTGC AGCTAAAGAT	10680
GTATTTGGTC TTTCTGATGT TGTTGCCTT GGTGGAACAA GTCTCTTGAT CATTATCTCT	10740
ACAGGTATCG AAGGAATCAA GCAATTGGAA GGTTACCTAT TGAAACGTAA GTATGTTGGT	10800
TTCATGGACA GAACAGAATA AAAGTATTAA CTGAATCAGT AAATACTGAG GGAGTCGAGG	10860
TTTAAACTCT GACATTGTA AGAGTTGGAT CTCCCCCTTT CTATTTGTT TTTAAATCGG	10920
GGTAAAAAGA CTTTTGCTT CTATTTAAAA ATAAAATAAG GAGATCAAAT CATGAATCTT	10980
TTGATTATGG GCTTACCTGG TGCAGGTAAG GGAACACTCAAG CAGCAAAAT CGTAGAACAA	11040
TTCCATGTTG CACATATCTC AACACGTGAT ATGTTCCCGC CTGCAATGGC AAATCAAAC	11100
GAAATGGGTG TTCTTGCTAA GTCATATATT GACAAGGGTG ATTGGTTCC TGACGAAGTT	11160
ACAAATGGAA TCGTAAAAGA ACGCCCTTCA CAAGATGATA TTAAAGAAAC AGGATTCTTA	11220
TTGGATGGTT ACCCACGTAC AATTGAACAA GCTCATGCCT TGGACAAAAC ATTGGCTGAA	11280
CTTGGCATTG AACTAGAAGG TGTTATCAAT ATTGAAGTGA ACCCTGACAG CCTTTGGAA	11340
CGTTTGAGTG GGCGTATCAT CCACCGCGTA ACTGGAGAAA CTTTCCACAA GGTCTTTAAC	11400
CCACCACTTG ACTATAAAGA AGAAGATTAC TACCAACGTG AAGATGATAA GCCTGAGACA	11460
GTAAAACGTC GTTTGGATGT TAATATTGCT CAAGGAGAAC CAATCATTGC TCACTACCCT	11520
GCCAAAGGTT TGGTCATGA CATCGAAGGT AATCAAGATA TCAATGATGT CTTCTCAGAT	11580
ATTGAAAAAG TATTGACAAA TTTGAAATAA AGCGTTTTTC ACACTTGCAA AAATCCGCTA	11640
CAAATGTTAT ACTGAGATAG TCTGACTTAT AATTGTTGTC TCTGTGTCTA GAGGCATCGA	11700
ATCGAAATTT ATGGAGGTGC TTTTGCCTGG CAAAAGACGA TGTGATTGAA GTTGAAGGCA	11760
AAGTACTTGA TACAATGCCG AATGCAATGT TTACGGTTGA ACTTGAAAT GGACATCAGA	11820
TTTTAGCAGG G	11831

(2) INFORMATION FOR SEQ ID NO: 66:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 10726 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double

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(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 66:

CCCGGCATTT GAAAGCTATT CGTGAAGGAT TTATGATGGC AATGCCTTG ATTTTAGTCG	60
GCTCTTATT TCTTATTCTA ATCAGTTGGC CTCAAGAGGC TTTTACAAAT TGGCTGAATA	120
GTGTTGGATT GCTAAGTATC TTGACAACTA TGAATCAGTC AACAGTAGCG ATTATCTCCT	180
TGGTCGCTTG TTTGGTATT GCCTACAGGT TGTCGGAAGG ATATGGTACA GATGGTCCGT	240
CGGCAGGGAT CATA GCCTTA TCCAGTTTG TATTGATGGC ACCTCGTTT TCGAGTATGG	300
TTTATGATAA AAATGGGGAG CAGGTCAAGC AGTTATTTGG CGGCGCAATA CCATTTCTA	360
GCCTGAATGC ATCTTCTTG TTTATGGCGA TTACTATTGG ATTGGTTACA GCAGAGATT	420
ATCGTATGTT TATCCAGCGC GGAATTACGA TAAAAATGCC AAGTGGTGTCCAGATGTAG	480
TAAGTAAATC ATTTTCAGCT CTTTTATCTG GTTTTACTAC TTTTGTGGT TGGGCTTTGG	540
TCTTAAAAGG TCTTGAAGCG GCAGGAGTTG CAGGAGGTCT CAACGGACTC CTAGGTGCAA	600
TTGTTGGAAC ACCGCTTAAG TTAATTGCAG GAACGCTTCC AGGTATGATT CTATGTGTTA	660
TTGTAAACTC ATTCTTTGG TTCTGTGGAG TTAATGGGG ACAAGTTTA AATGCTTTG	720
TAGACCCAGT TTGGTTACAA TTACTACAG AAAACCAAGA AGCTGTGGCT GCAGGACAAA	780
CACTCCAACA CATTATTACA TTACCGTTA AAGATTTATT TGTATTTATT GGTGGCGGTG	840
GAGCGACTAT TGGTCTTGCAT ATTGTCTCT TCCTATTTAG TAAGAGTCGT GCGAATAAAA	900
CATTTAGGTA GCTAGCTATT ATACCGCTA TTTTTAATAT CAATACAGCT ATTCTATTAA	960
CGTTTCCAAC AGTTTTAAAT CCGATTATGC TGATTCCGTT TATTGCTACT CCTACAATCA	1020
ATGCCTTGAT TACCTATGTA TCAATGGCTG TAGGATTAGT ACCCTATACA ACAGGTGTAA	1080
TCCCTCCGTG GACAATGCCA CCGATTATAG GAGGCTTCCCT TGCAACAGGG GCTAGTTGGC	1140
GAGGAGCTCT ATTACAAGTT GTTTGATTT TGTTTCTGT AGCAATTAT TATCCATTCT	1200
TCAAAATTGC AGATAAACGC AATCTTGGAA AAGAAAAAGC TACTGTGGA GGGAAATAAG	1260
ATGGTTATCA GAGTATTGTA TCAACAGAAA AATACTTATT CTAGCTTGC CTTAGAGGAA	1320
TTAAGTTACT ATATGAATCG GGTCTTTAAG ACTAACATAG AGCTGTGCGA GGAGAAGGAA	1380
CGGGATATTG TTGTAGGATT AGTCAATAAA GAGGACAGAA AAGACCAGT TCTTATCTCA	1440
TTAGACAAGG GTAAGGGGAG AATTGAGTCT AATACAATTG TAGGTTACT TATTGGAATT	1500
TACCGAATGT TTCATGAATT TGGGGTTGTG TATACTAGAC CAGGGCGCAG ACATGACTTT	1560
GTTCCAGAGT TACGATTGTA AGATTTTTA GATAAACAGC TATCTATAGA TGAAACAGCC	1620

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AGTTACTATC ATAGGGAGT ATGTATAGAG GGAGCGGATT CATTGAAAAA TATACTAGAT	1680
TTCATTGATT GGCTACCTAA GATTGGGATG AACAGTTTT TCATCCAGTT TGAAAATCCT	1740
TACTCTTTTG TGAAACGTTG GTATGAACAT GAATTAAATC CATATCTAAA TAAAGAACAA	1800
TTTTCAAAATG AATTAGTACA AGAATTGAGT GATAGGTTGG ATAAAGAATT GCAAAAAAGA	1860
GGTCTTATTC ATCATCGTGT TGGTCATGGA TGGACAGGTG AAGTTTTAGG TTACTCTTC	1920
AAATTTGGCT GGGAAATCAGG TCTTAGTATT TCAGAGGAGA AGAAACCTA TGTCGCTGAA	1980
ATAAACGGGA AACGAGAATT GTTTAATACG GCTCCGATTT TAACCAGCCT GGATTTTPCA	2040
AATCCAGATG TAGCTGATAA GATGGTAGAA ATTATCAAGG ATTATGCCAA GAAAAGACCT	2100
GATGTTAACT ACTTACATGT ATGGTTGTCG GATGCTCGTA ATAATATTTG TGAATGCCAA	2160
AACTGTAGAC AAGAATTGGT TTCCGGATCAG TATATTCGTA TTCTCAATCA ATTGGATAGG	2220
GCTTTAACGA GTGAGGGATT AGATACAAAG ATTGTTTTTC TGCTTTATCA TGAGTTGTTA	2280
TGGGCACCTC AGAAAGAAAA ATTAGATAAT CCTGAACGCT TTACCATGAT GTTTGCACCG	2340
ATTACAAGAA CATTGAAAT GAGTTATGCA GATGTAGATT TTGACAATTC CATAACCTACG	2400
CCTAAACCTT ATATGCGTAA TAAAATTATA CTTCCGAATT CTCTTGAGGA AAATTTATCT	2460
TATCTTTTG AGTGGCAAAA AGCATTAAA GGAGATAGTT TCGTATATGA CTATCCTTTA	2520
GGGCGTGCCTC ATTATGGCGA TTTAGGCTAT ATGAAAATTAA GTCAAACCTAT TTACAGAGAT	2580
GTATCTTATC TTTCACACCT ACATTTGAAC GGGTACATT CGTGTCAAGA ATTACGTGCC	2640
GGATTCCCTC ATAATTTCC TAATTATGTC ATGGGGAAA TGCTCTGGAA GAAGACAAGA	2700
AGTTATGAAG AATTGATTGA AGAATACATT TCTGCTTTGT ATGGGGAAAA TTGGCAGTCT	2760
GTTGTTGAAT ATTTAGAAAA ATTATCCATT TATTCTCTT GTGATTATTT TAATGCAATT	2820
GGCAGCGTC AAAGTGTGTT TTTAGCGAAT CATTATTATA TAGCTTACAA TCTAGCTGAT	2880
AATTTTTAC CAATTATTGA GGAAAATATT TCTAAGTTAT TAAATAGTCA AAAGGATGAA	2940
TGGAAACAGC TCAGTTATCA TCGTGAATAT GTGTTAAGA TGGCGAAGGC TTTATATCTT	3000
CAAGCCAATG GAAAACAAG GCAAGCTCAA GATGAATGGA GAAATGTGTT GAATTATATC	3060
CGTGGGCACG AATTGCTATT TCAATCTAAT TTGGATGTTT ATCGTGTAAAT TGAAGTAGCA	3120
AAAAATTACG CTGGTTCCA CTTATAATC ATAAGTATAG AAAATGAACT AAGGTATTCA	3180
GAGAAGATTG ATCCTAAATA TTATGAAATT TAAGGATTTT TAAGATATTG AGGGTCAACT	3240
TTCTATTTAT ATCGTAGCGA AGTCATTTA ATAATGATGT GTAAAAGATG GATCAAGATT	3300
GAGGAGGAAG AAAGATGAAA TCAAAAGAAG AAATAATAT GCTTGGTTTT ACAATTGTCG	3360

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CTTACGCAGG AGATGCAAGG TCAGATTG TGATGCTTT GGCGTTGCG AGAGATGGAT	3420
ATTTTGAAACA GGCAAGAGAA TTGGTTGAGT CTGCAAACGA CTCATAAGTG TCTGCCCATC	3480
GAGAACAGAC TAATTATTA GCGGAGGAGG CATATGGAGA TAATTTGAA GTGAGCTTTA	3540
TTATGATTCA TGGTCAAGAT ACTTTGATGA CAACGATGCT ATTGTATGAT CAGGTAAAGT	3600
TTTTTATTGA TGAATATGAA CGAATTGAA AGATTGAAGA ACATATTGGT TTGCAATGAG	3660
GATTAGTCAT GGAAAATTAA CAGGTTAAAG CCTTACCGAA GGAGTTTTA TTAGGAACTG	3720
CTACCGCTGC TTATCAAGTA GAGGGTGCAA CTAGGGTAGA TGGCAAAGGA ATAAATATGTT	3780
GGGATGTTTA TTTGCAAGAA AATAGTCGCT TCTTACCGAGA TCCAGCTAGT GATTTTATT	3840
ATCGTTACGA AGAGGATATA GCTTGGCGG CAGAACATGG TTTGCAGGCT TTGCGTTTAT	3900
CTATTTCTTG GGTCGTATA TTTCCGTATA TAGATGGGA TGCTAATGTA TTAGCTGTTTC	3960
ATTATTACCA TAGAGTTTT CACTCTTGCT TAAACATAA TGTGATTCCG TTTGTTTCTT	4020
TACATCATTG TGATTCGCT CAGAAAAATGT TAGAACAGG GGATTGGTTG AACAGAGAGA	4080
ATATTGATCG TTTCATACGA TATGCTCGCT TTTGTTCCA AGAATTACAA GAAGTCAGC	4140
ATTGGTTTAC AATCAATGAA CTGATGTCTC TTGCTGCAGG TCAATATATAA GGAGGTCAGT	4200
TTCCCTCCAAA TCATCATTG CAATTATCTG AAGCAATTCA AGCGAATCAT AATATGTTGT	4260
TGGCGCATGC TCTTGCAGTC CTCGAATTTC ATCAATTAGG GATTGGGA AAGGTAGGTT	4320
GTATTTCATGC TTAAAGCCA GGCTATCCTA TTGATGGCA AAAAGAAAAT ATTTGGCAG	4380
CTAAACGGTA TGATGTTAT AATAATAAT TTCTTATTAGA TGGAACTTTT TTGGGCTACT	4440
ACAGTGAGGA CACCCCTTTT CACTGAATC AAATATTGGA AGCTAATAAT TCTAGCTTTA	4500
TTATTGAAGA TGGTGTGTTA GAAATTATGA AGAGAGCTGC ACCTCTTAAT ACGATGTTG	4560
GGATGAATTAA TTATCGTTCA GAATTATTC GTGAATACAA AGGTGAAAAT AGACAAGAAT	4620
TTAACCAAC AGGAATAAAA GGACAGTCTT CTTTAAATT AAATGCTCTA GGTGAATTG	4680
TAACACACCC TGGTATTCCG ACAACAGATT GGGATTGGAA TATTTATCCT CAAGGGTTAT	4740
TTGATATGTT GCTTCGTATC AAAGAAGAAT ATCCTCAACA TCCGGTCATT TATTTAACTG	4800
AAAATGGTAC AGCCCTTAA GAAGTTAACG CAGAGGGCGA GAATGATATT ATTGATGACA	4860
GTAAGAGAAT CCGTTATATT GAGCAACATT TACACAAAGT TTGAGGCT CGAGATAGAG	4920
GAGTCATAT TCAAGGCTAT TTATATGTT CTTTGCAGA TCAATTCTCT TGGCGAATG	4980
GCTACAATAA GCGATATGGT CTTTCTTTG TTGATTATGA AACACAGAAG AGATATATTA	5040
AGAAAAGTGC TCTTGGGTA AAAGGGCTAA AACGGAATTA AGGTTAGCGA TTTGACTGAT	5100
GTTTAATATG TTTAAATAT GAGGTGAAT TTTTATAGG AGGAGTTTTA TGGATAAGCT	5160

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AGTCGCTGCC	ATTGAAAAGC	AACAAGGGAA	ATTTGAAAAA	ATTTCTACTA	ATAACTATAT	5220
GATGGCTATT	AAAGATGGAT	TCATTGCTAC	TATGCCTTA	ATTATGTTT	CAAGCTTTT	5280
GATGATTATT	ATTATGATTCT	CTAAAAATTT	CGGAGTAGAG	TTACCGAGTC	CAGCTATTGT	5340
CTGGATGAGA	AAAGTGTATA	TGTTAACCAT	GGGAGTTTG	GGTATTATTG	TTTCAGGGAC	5400
TGTTGAAAG	TCATTAGTTG	GAAATGTTAA	CAGAAAATG	CCTCACGGAA	AGGTAATAAA	5460
TGATATTTCT	GCAATGTTGG	CAGCCATATG	TAGTTATCTG	GTATTAACTG	TAACGCTTGT	5520
AGTTGATGAG	AAGACGGGAT	CTACAAGTTT	GTCGACAAAC	TATTTAGGAT	CTCAAGGATT	5580
GATAACTTCG	TTTGTCACTG	CCTTTATTAC	TGAAATGTT	TACCGATTCT	GTATTAAGCG	5640
AGACATTACT	ATTCATTTAC	CTAACAGGAAGT	TCCTGGGGCT	ATATCACAAG	CTTTTAGAGA	5700
TATTTTCCCT	TTTCTTTTG	TTTTACTTAT	TAGTGGTTTG	TTAGATATTG	TATCTCGGTT	5760
TAGTTTAGAT	GTTCTTTTG	CCCAAGTATT	TCAACAACTA	TTGACTCCTA	TTTTTAAGGG	5820
GGCAGAACATCA	TATCCTGCTA	TGATGTTGAT	TTGGTTTATG	TGTGCTTTGC	TTGGTTTTGT	5880
TGGAATTCTAT	GGACCATCTA	TTGCTTTACC	TGCTGTTACA	GCTTGCAC	TGAGCAATAT	5940
GGAAGAGAAAT	GCTCAACTTC	TTGCAAATGG	GCAGTTCCCT	TATCATTCTT	TAACACCTAA	6000
TTTCGGAAT	TATATCGCTG	CTATTGGAGG	AACGGGGGCT	ACCTTGTTG	TACCATTAT	6060
TTTGATTTTC	TTTATGCGGT	CTAACACAATT	AAAATCGGTA	GGTAAAGCTA	CAATTACTCC	6120
TGTTTTATTT	GCGTAAATG	AACCTCTTCT	ATTGGTATG	CCTGTTATTT	TGAATCCCTA	6180
TCTTTTTGTC	CCTTTTTGTA	TGACTCCACC	AGTGAATGTA	TTTCTAGGAA	AGGTCTTTAT	6240
TGATTTCCTT	GGAATGAATG	GATTTATAT	CCAGTTACCT	TGGACCTTTC	CTGGTCCCTT	6300
GGGATTGTTA	ATTGGAACGA	ATTTTCACT	TATCTCCTT	GTATTTTAT	CTTTGATTTT	6360
AGTTGTCAC	ATATTGATTT	ATTPGCCATT	CTGTAGAGCG	TATGATAGAC	AGTTACTGGT	6420
GAAAGAAGAT	ATTGCAAGCT	CAAATGATAT	TATTTTAGAG	GAGGATACAA	GTGAAATAAT	6480
TCCTGGTGAG	ATAGATGAAA	AAAAAGTAA	GGAGTTGAAA	GTACTGGTTC	TTTGTGCAGG	6540
GTCTGGAACA	AGTGCACAA	TAGCCAATGC	AATTAACGAG	GGGGCTAACT	TAACAGAGGT	6600
TAGAGTGATT	GCGAATTTCAG	GAGCGTACGG	AGCTCATTAT	GATATTATGG	GTGTTTATGA	6660
TTTAATTATT	CTGGCCCCAC	AAGTCGGAG	TTATTATAGA	GAGATGAAGG	TGGATGCAGA	6720
AAGATTAGGT	ATTCAGATAG	TTGCTACCA	AGGAATGGAA	TATATTCA	TAACAAAGAG	6780
TCCAAGTAA	GCCTTACAAT	TTGTATTGGA	GCATTACCAA	GCTGTAGT	AAGTTTTCC	6840
ATCTTTTATT	TGAGTAAAGA	TTTGTTTAC	AGATAGGCTT	GGATTTAAAA	ACGTTCCCCC	6900

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TTTTTTAATA TAAGAACCCC TCTTTCACAA TTGTAAAAAG AGGGATTTG TATTTTATCT	6960
CTTAGACCAA GTTCTCTTCA TAAAGAGAAG GAGGATTGGG TAAATCTCCA AGCGCCCTGC	7020
AATCATTGCA AAGGATAGGA GAATTTTGAGA GATGGGACTA AAGATTGAGA AACTAGAAGT	7080
GGTTCCCTAGA ATAGGCCGA TATTATTGAA ACAGCTAAAG ACAGCGCTGG TCACGACCAG	7140
AAAATCATTG CTATCTAGGC TGACAATAAA GATAAGCGCT AGCAAAATCA TAGCATAGAT	7200
GACAAAGTAC TTGAGAATCT TATGCTGGGT ATCTTGTCA ATCACCGTTT TATTAACATG	7260
GAGGGTCAAA ACACGGTGGG GCGATAGGAT TGACAAAATT TGGTTTTGG CAATTTTGAGA	7320
AAGGATGAGG CCTCGAATAA TCTTGAGTCC ACCTGCAGTT GATCCAGCAG AGCCACCGAT	7380
TGCCATGAGG AAAAGGAGGA TAAACTGGGA GAAGAGGGC CAGTPGGTAA TATCTCCATA	7440
TCCAAAACCA GTTGTGTAA TGATGTTGGA AACCTGGAAG AAGGTCATTT CAAAGCTCTT	7500
TGAAAACCTT GGCTAGAGGT AGAGGGTGTGTT GAGGCTAATC AAGCCTGTAG AAACCACTAC	7560
AATGACCAAG TAAGCCCTAA GCTCTTCATC TCCAAAGAAG GCCTPGATGC GACGGAGCAT	7620
GAGGTAGTAG TAGAGGTTGA AATTTACTCC AAAAACAGA ACTCCGATAC TGACCAGATA	7680
GGTAATCAGT GAGCTGCCAT AGTGGGCAAT TCCGTCGTTA TAGACGGTAA AGCCTCCAGT	7740
TCCCGCTGTC CCCATAGCAA TAACAAAATC ATCGTAGAGA GGCATACCGG CTAGATAATA	7800
GATGATGACA AAGAGGGAGA AGAGAGCTAG ATAAAGGAGA TAGAGAATCT GGGCAGTGT	7860
TTTTAGTTG GATACAACCT TGCCAAAAG AGGACCTGGA ACCTCAGCCT TCATCACCTC	7920
TAGGTGGCTA TTTTTGGCAT TGTCATAAT AGCAAGTGC AAAACAAGCA CTCCCATCCC	7980
TCCAATCAAG TGGTAAAAC TTCCGCCAGAA GAGGAGGGAA CGGCTGAGAA CCGAAACGTC	8040
GTTCAAAAATA CTTGCTCCAG TAGTTGTAAA TCCAGAACTA ATTTCAAAAA AGGCATCAAT	8100
AAGGCTGGGG ATTTGCCAG AAAAGACAAA GGGGAGACCA CCAAAGAAAG ACCAAAGGAT	8160
CCAAACAGAGG GCAACGATCA AGACTCCCTC CTTGGCATAA ATCCGTTGAT TTTTTGGCTT	8220
CTGTAAACTC CCTGAACCGC CTAACAATAC GAGAATCCCT ATGGTCGAAA AGAGGGCTGT	8280
AAAGACTTGG CTCGATTACAG GGTAAATAGAC AGCAATCGCA ACAGGAACCA AAAGAAGAAC	8340
AGCTTCAATC AAAAGTAATT TTGAAAGGAG GTAACGAATC ATACTTTAT TCATTTCTTA	8400
CCTCGCGATC AAGTCATAAA TCTTGGTGAT GTTTGGCAAC AAGGTTGTTA CTAGGAGCTT	8460
GTCTCCAATC TCCAACATAT CCTCCCCAGT TGGGAAAATA GTCTTGGCCTT TTGCAATAAT	8520
GGCTGCAATA AGAACCCCTT TTTCAATT CAGTTGAGAA AGAGGTTGG CAGTCATT	8580
ATTGGCTTCC TTGATATGGA ATTGCAGGGT TTGATTTGG CCATTGGCTA GATGGTGCAT	8640
AGCTTGAAGG TCTGAATACT GGGCATTAAC TCGACCACGA ATAAAGTGCA TAATCGTATC	8700

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TACAGCGATG	CTTTAGGTG	TGATGATACT	TGAAAATCA	GGCGCATTGA	TAATCTCGAG	8760	
GAGACTGGTA	CGATTGACCT	TAGTAATATT	TTTCTGTACA	CCTACCCCTGT	CAAGGAACAT	8820	
AGATGTAATC	AGATTTCCCT	CATCGACTCC	TGTTAGAGTC	GCAACGGCAT	CATAGTGTG	8880	
AGCACTTTCT	TCCAGCAGGA	TATCTTTGC	GGTTCCATCT	CCTTGAACGA	TGTAGAGATT	8940	
TGGGAATTTTC	TCGCTAAAGA	AGCTGGCGAT	TTCAGGATTG	ATTCATGA	CTTTGTATC	9000	
GATACGACTA	TCTTGAGAA	TACCAAGTAG	ATAATAGGCA	ATTCTACCTG	CCCCAACGAT	9060	
GAGAAGGCTC	TTCACGGCGC	GTGATTAAA	ATAATTATGG	AAGAGTATCA	TATCGACACG	9120	
GTTACCAGTG	ACAAGATTTC	TATCTTTATC	CTGTACAGTC	ATGTCACCGC	TTGGAATGAT	9180	
AATTTGATGA	TCCCCTCTCTA	TCGCACAGAC	AATGACATTA	CCAAATTTT	TACGAAAATC	9240	
AGAAATGGGC	ATTTGGCAAA	GACCGCTGGT	GGACTTGACG	ACAAATTCCA	TGAGGCTAAC	9300	
GCGTCCACCA	GCAAAGCGTT	CGACAGACAG	GGCGTTGGGG	AAGTCATGA	TATTGCGCAT	9360	
AGCGCGGGCA	GCCAAGAGCT	CAGGATTAAC	GATAAGAGAA	AAACCGAGAA	TATTCTTTTC	9420	
CTTGAATAA	GAGTTAGAAT	ATTCAGGGTT	CCGCACCCGA	ACGATAGTTT	CTTTAGCTCC	9480	
CATTTCTTG	GCTAGAACTG	CTGCAATCAT	GTTGACTTCA	TCGTGCTCAG	TCAGGGCGAT	9540	
AAAGATATCA	CAATCTTGGA	CGCTGGCTTG	CTCAAGAATG	GCAAATCGG	CCCCGTTACC	9600	
AAGGATACCA	ATGATATCAA	AGCGACTGAC	AAATGATTG	AGAACAGCTT	CGTCCTGCTC	9660	
AATCAGCAAA	ACATCATGCT	TTTCTGCAAC	CAAGGAGCGA	CAGAGGGCAA	AACCAACTTT	9720	
TCCCCCTCCG	ACAAGGATAA	TTTCATAAT	AAAACCTACT	TTTCATGAT	GTAACATATCA	9780	
TACCCCTTTT	CAAGAAAAAA	TGCACCTACT	AGCTAATAAC	AAGAGTTTT	AGTGAAAATT	9840	
CGCTATAAGG	TTAAACTATA	CCCTAACCAA	TTGAAATAGC	TATTAGCGAC	TTTCTCTGAA	9900	
ATATGGTATG	ATAAAGGATA	TACAAGGAGA	TTAAATGAAT	ATAAATTTCAC	TGGTATTACA	9960	
ATCAGACTTT	GGTCTGGTTG	ATGGTCCGGT	ATCGGCTATG	ATTGGAGTGG	CTTTAGAAGA	10020	
GTCTCCAACC	TTAAAAATAC	ATCACTTGAC	GCACGATATC	ACGCCCTATA	ATATTTTGAT	10080	
GGGGAGCTAT	CGTCTCTTTC	AGACGGTGGA	TTACTGGCCT	GAGGGAACGA	CGTTTGTATC	10140	
GGTTGTCGAT	CCAGGGTGTG	GTTGAAACG	TAAGAGTGT	GTTGCCAAGA	CTGCAAAAAA	10200	
TCAATACATT	GTCACGCCAG	ATAATGGGAC	GCTTCCCTT	ATCAAGAAC	ACGTTGGCAT	10260	
TGTAGCCATT	CGTGAGATT	CTGAGGTTGC	CAATAGGCGT	AAAACACAG	AGCATTCTTA	10320	
TACCTTCCAC	GGTCGTGATG	TCTATGCCTA	TACTGGTGCT	AAACTGGCCA	GTGGTCACAT	10380	
TA	CTTTGAG	GAAGTAGGGC	CAGAGCTCAG	TGTGGAACAG	ATTGTAGAGC	TTCCAGTCGT	10440

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AGCGACCATC ATAGAAGATC ATCTGGTGAA GGGAGCCATT GATATTCTGG ATGTGCGTTT	10500
CGGTTCGCTT TGGACCTCTA TCACACGGGA AGAATTTAC AAGCTGGAAC CAGAATTG	10560
TGATCGTTT GAAGTGACCA TCTATCATGC TGATATGCTG GTCTATCAAATCAGGTTGT	10620
CTATGCCAA TCATTTGCAG ATGTGAGAAT TGGGCAACCS ATcTTTACrc TCAGCaTCTt	10680
CGATTAGCTG GGCAATTCTG TCTAGTTGGA TTTCGTCAAT CAAGGT	10726

(2) INFORMATION FOR SEQ ID NO: 67:

- (i) SEQUENCE CHARACTERISTICS:
- (A) LENGTH: 7163 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 67:

TTATCTTAA CGATATCAAT CAAGATCTGG TCAATAAAGG GATTGGGCT TATCGTGAAG	60
TTGGCATCCA AGCCCATGGA TATGTCTGTG ACGTGACAGA CGAGGACGGT ATCCAAGCCA	120
TGGTCAAGCA AATCGAACAA GAGGTTGGTG TCATTGACAT CCTCGTTAAT AACGCTGGTA	180
TTATCCGCCG AGTTCCAATG TCCGAAATGA GCGCCGCTGA TTTCCGTAAG GTCATCGATA	240
TTGACTTAAA CGCACCAATT ATCGTTCAA AGGCAGTTAT TCCTTCTATG ATAAAGAAAG	300
GGCATGGAAA GATTATCAAT ATTTGTTCGA TGATGAGCGA ACTGGGACGT GAAACAGTTA	360
GCGCTTATGC TGCTGCTAAA GGGGGCTTGA AAATGTTGAC CCGCAACATT GCGTCTGAAT	420
ACGGTGGAGC CAATATCAA TCTAACGGAA TTGGACCGGG TTATATTGCC ACTCCTCAA	480
CAGCACCTCT TCGTGAATTG CAAGAAGATG GTTCTCGCCA CCCATTGAC CAGTTCATCA	540
TTGCAAAAC ACCTGCTGCA CGTTGGGAA ATACTGAAGA TTTGATGGC CCTGCTGTCT	600
TTCTCGCTAG TGATGCCAGC AATTTCGTCA ATGCCACAT CCTATATGTA GATGGCGGTA	660
TCTTAGCCTA CATCGGAAA CAACCTGAGT AAAAATAGAA AGAAGATCTT ATGAAAATCG	720
CATTAATCAA TGAAAATAGT CAAGCTAGCA AGAACATCAC TATTTACGAT AGTCTAAAAG	780
AAGCGACAGA TAAAAAAGGC TACCAATTAT TTAACATATGG TATGCGTGGAA GAAGAAGGAG	840
AAAGTCATT AACTTATGTG CAGAACGGAC TAATGGCTGC CATCCTTTA AATACAAAGG	900
CAGTTGACTT TGTTGTTACG GGTGTGGTA CGGGTGTAGG GGCTATGCTT GCTTAAACA	960
GCTTCCTGG TGTTGTCGTG GGTCTAGCAG TGGACCCAAAC TGACGCTTAC CTTTATTCTC	1020
AAATCAATGG TGGTAACGCC TTGTCTATCC CTTATGCCAA AGGATTGGC TGGGGGGCAG	1080
AACTGACCCCT CAAATTGATG TTTGAACGCT TATTTGCTGA AGAAATGGC GGTGGCTACC	1140

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CAAGAGAACG TGTAATCCCT GAAACAACGCA ACGCTCGTAT CTTAAACGAG GTGAAACAAA	1200
TCACCCACAA TGATTTGATG ACCATCCTTA AAATAATCGA CCAAGACTTC CTCAAAGACA	1260
CCATCTCTGG CAAATACTTC CAAGAATACT TCTTGAAAA CTGCCAAGAT GATGAAGTTG	1320
CTGCTTATTT GAAAGAAGTA TTAGCCAAGT AAAGCTATTG TAAACCAGAA AGGAACATAAT	1380
GGATGACGAA AATATTACTG TTTGGCGAAC CATTAATTG AATTTCACCA TTAGATGCCA	1440
CCAGTATCGG CGATCATGTT GCCAGTTCGA CTTATTTGG CGGATCAGAA ATTAACATCG	1500
CTTGTAATTT GCAAGCCCTG GGTATCTCAA CGAAAGTTT TACCGCACTC CCTGCCAACG	1560
AGATTTGGAGA TCGTTTTCTC ACATTCITGA AACAGCACCA AATCGATACC AGTTCAATCT	1620
GTCGGCTTGG CGATCGAACG GGCCTCTACT ATTTGGAGAA CGGCTTGGT TGTCGTCAA	1680
GTGAAGTTT CTACGATCGT AAGCATAACGA GTATCAGCCA GATTCGGCCA AACATGCTAG	1740
ATATGGATTC TCTCTTTCAAG GGGATTAGCC ATTTTCATTT TAGTGGAAATC ACCGTAGCTA	1800
TCGGTCAAGA GGTCCGTGCG ATCCTTCTCC TACTCTTGGG AGAAGCCAAG CGCCGAGGAA	1860
TTGTCGTTTC AATGGATCTC AATCTGAGAA CAAAGATGAT TTCAGTCCTA GAAGCCAAGT	1920
ATGAATTTTC TAAGTTGCA CGTTTTACTG ACTATTGCTT CGGTATTGAT CCTCTCATGA	1980
TTGATGACCA AAATCTAGAG ATGTTTCCAA GAGACAGTGC TAGCCTAGAA GAGGTGGAAA	2040
ATCGCATGCCG ACTTTTAAAAA GAAGCCTATG GTTCAAGGC CATTTCCTCAT ACCCTCCGCT	2100
CTACTCMTGA GCAAGACAAA AATGTCTATC AAGCCTATGC TCTAGAAGAA CTATTTGAAG	2160
AGTCTGTCCA ACTAAAAACT GCAGTCTATC AACGAATTGG TAGCGGGGAT GCCTTTATAT	2220
CTGGTGCCTT TTACCAACTA CTCCATCATT CCTCCCTAAA AACTACCATT GACTTTGCAG	2280
TTGCGAGCGC AACTCTCAA TGCACTCTTC CAGGAGACCA TCTCTCCACT TCCTCAACTA	2340
GTATTGAAAA TTACTGGCA AATGCACAAG ATATCATTG TTAGGAGAAAT TACATGACCA	2400
AATCAGATAC GATTATTGAA CTAAAAAAAC AAAAATTGT CGCTGTTATT CGAGGAAATA	2460
CAAAGGAAGA AGGACTACAA GCCTCGATTG CTTGTATCAA GGGCGGTATC AAAGCTATTG	2520
AAATCGCCCTA TACCAATCAG TATGCAGGAC AAATCATCAA GGAACCTGTA GACTTGTATC	2580
AGGACGATCA GAGTGTGTTGT ATCGGTGCGAG GTACTGTGCT TGATGCCGTA ACTGCTAGAG	2640
ATGCCATTCT AGCTGGAGCA AATTACGTTG TTTCTCCATC TTTCCATGCT GAAACTGCGA	2700
AAATGTGCAA TCTCTACAGC ACACCGTACA TTCCAGGCTG TATTACCCCTC ACAGAGATCA	2760
CGACTGCCTG TGAAGCCGGT AGTGAATCA TCAAACCTTT CCCAGGTAGT ACTCTCAGTC	2820
CAGCATATAT CTCTGCAGTC AAGGCACCGA TCCCACAAAGT TTCCGTAATG GTAACCGGAG	2880

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GAGTCGGCCT	AAACAACATC	CCTCAATGGT	TGGCTGCTGG	TGCAGATGCC	GTTGGAATTG	2940
GTGGCGAACT	CAATAAACTC	GCTTCCAAG	GCAACTTTGA	CCGCATCAGC	GAGATTGCC	3000
AACAGTATAT	TACACTCAGA	AAAATCATA	ACTACCCGTC	TAACGGGTGG	TTTATCTCAG	3060
AGCTATAAGC	CCAAATCATC	AGCCAGCGCC	TAAAGACGCT	GGCTTCACG	TTGTTCAAGC	3120
CTTATTGCTC	TTGACTCGTC	ACTTGCTCT	TTAAGAGACT	TTGGTATTAC	TTACCACTAT	3180
CCCTAAAGGG	ATCCTCATAT	TCTTTACAC	TCAATTATC	TAGTGTATA	GTAGATTGAA	3240
ACTGGAATAG	TACACCTCTG	CTTCTAAAAC	ATTGTTAAA	ATCGATTGAA	CTGTCCTGAT	3300
CGATTTGTC	CTGTTCTTAT	TTCATTTAC	TATATATCAT	ACTTTACTCG	TTCTCAAATT	3360
TTCATACTCA	TGAAGAAATC	ATCCACTCGA	TAATTCTTT	AATCTTGA	CTATTTCTTA	3420
ATTGTGGCTT	CATTAAGCCC	TACTGGACTT	ACATAATAAC	CTTCCTCCCA	GAAATGCCGA	3480
TTCCCCAACT	TGTACTTGTG	ATTGGCGTGT	TTGTCAAACA	TCATGAGTGC	ACTTTGCCT	3540
TTTAAATACC	CCATAAAACT	TGAAACACTT	AGCCTCGACG	GAATACTGAC	TAACATGTGT	3600
ACATGGCTG	GCATTAAGTG	ACCCCTGATC	ATTTCAACAC	CTTTATAACT	ACACAAGCGA	3660
TGAAATATTT	CGTCTAAACT	ACTTCTATAT	TGATTATAGA	TGACTTTTCG	TCTATACTTA	3720
GGGGTGAACA	CAATATGATA	GAACACCTCC	ACTTTGTGTA	TGATAAACTA	TGAGTCTTT	3780
GTGCCATATT	TTTCTCCTT	TCGCTTACA	ATTGGATTGA	ACACCTTTAT	TGTATCGCGT	3840
TTGGAGTTT	TTTGGTATAA	CCTCGACGC	GCACCCGTAT	AGCGGGTGGT	TGTTTTGTCT	3900
CGCACCTCAC	GGAGGGAGAC	GGACTAATAT	AGTGGAGTGA	AATAGGATAC	GAACAAATTG	3960
ATTAGGAAAA	TCAAATGAAT	TTATAGAAAT	CTTTTAGCAG	TTATAACGTT	CTATTCTAGT	4020
TTCAAAACGC	TATAGTCACA	TAATAATGAA	GTAAAAAAGG	ATAAGTATCA	ACTTATCCTT	4080
TTTTAAAAAGA	AAAATCCGAA	GATATTTGGC	CTTCTTCGGA	TTTTTTCTAT	TTTCCACAGT	4140
TTCATGTAAT	TCATCTAGAT	GATGAACAAA	TTAGTTGTC	TTTCCTCTAC	CCAATAGATA	4200
AAATGCCCA	AGTAGCAAGA	ACCCCTAGACT	TGCCAAGATT	GACTGACCTT	CTCCTGTCTG	4260
AGGGAGATTC	TTTGATCCG	AATGGTTCTT	TTCCCTTTCA	GATTTTTCCCT	TTTCTTTGA	4320
ATTCTGTA	TGTGGCTGAG	CTGCTTGCTC	TAGCTTTTA	AAGACTTCCT	GATCTGGAGC	4380
TGATTCCTGG	GTTCAGGAT	TATAGTAGGC	AATCTTATAT	TCATCCCTT	CTTTTCGAAT	4440
GGTATAGACT	CCACGTTCA	AAACTTGAA	TTGGTTGGAA	ATAGTAGAGA	CAGAACATC	4500
ATATTCACA	ATGCCCAAA	CTCCCTGTTT	AGCATCATAA	ACAGACTGAA	GGGTTTCGTT	4560
ATTTTCGATG	AGGCTACTTT	CTAACTCTT	TATCATTGAA	TTGAAGGTGG	CACGATCCAC	4620
GTTAGGAATG	AGCATATAGC	CATAAGAAC	TCTATTTGC	TTATGAGCCT	GACTAACCGT	4680

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AAGAAATTCA	TTTCAACTT	CCTGTCTGA	CTGTCCTTC	TTGATATCCT	TCCAGGCTCC	4740
CTTTTGCAAA	GCCTTA	CTCA	TACTGATTGA	ACTCTTCTTA	AAGAAAAAGT	4800
CTTTTCGAA	TCGAACGATT	CTAAAAAGAC	ACTTTGGGTT	TCAGGATAAT	CCTTTCTTG	4860
TTCTGTAAGG	GAGGCTCTT	TATCATTGAC	ATAGACTTTA	TATGGATTAC	CTGATTCCAG	4920
TTTTCTCTGG	TCAATTGAG	TTGAGCAGT	ATCTGTTGAA	GTGTTTGGG	TATTGCTTCC	4980
TAAAAAGGCG	ATCTTATCCT	TTAGCATAAA	CCAGCTCTTA	TGAGCAGTCA	ATGTTTGATT	5040
CCAGTTGGTG	AAATCCATGG	TTGCTGTCGC	ATTGGCATCA	TCTAGTTGCC	TCGTTCCAAC	5100
GAAAGCAGAC	GGTAAAACCTT	TACCTGTATC	GCTATCCGCT	CTCTTAGCAT	CCGCTCTGT	5160
TGTACCAGGC	ATCTTATATG	GATTAACGT	TGGCCAGTAG	CCATCGCTAT	AGTGACTCAA	5220
ATCGCCATTG	TAAAGATAGA	ACATCCCATC	ACTCGTATAC	CAACCACGTT	TATTTCCCTT	5280
GTTCATGTGT	TCGTAATTCA	AGGTACGACT	GGAAAAGAGT	GACAAGCCAA	ATCCAAACCC	5340
TTTCTCTGCA	TTGTACATGG	CTGTTTATC	CATCTGTTA	AAGGCAGATA	GGTAACTTGG	5400
TCTTGGAAACA	CTTGGCAGTC	CTGCATCACT	TAACAAGGAT	TGCATCAAAC	TGATATCCTT	5460
ATAAGTCTTC	AAATTCTAA	AGACATCATA	ATAACTATCC	GATTGAACAA	TGGTCTTCAC	5520
AAGACTCTGC	AAACATTGTT	TGGTTCTCC	TTCAGACATA	TCCGCTATTC	GGTGAATCCC	5580
TCTTAGTACT	TCTACTGCGG	CCACGTGCC	CTCGCTATTT	GCACGACTGA	TCGAGCGTCC	5640
ACGACTCATA	TCCATCAACT	CTCCATTAC	CAGCAAAGGA	GCAACACGATT	TATCAATCCA	5700
GTGGTACATG	GTTCGATTT	TATCTTATC	GATTGGATTC	TTGGTCTTTT	GAATGACTGG	5760
CAACAGTTGA	GACAGGCCAT	CAATCAAAAC	ATTCCCATAA	GCACCCGTAT	AGGCAACATT	5820
GGTGTGGTCG	ATATAGGATC	CATCTTGATA	AAAACCTTCA	CCTTGGTCTA	CCAACATTGAA	5880
CACTTGCTCA	ATCGAGCGAA	TGGTAGAAGA	AATTCTGTA	TCATCCTTAC	GCAGTAAACC	5940
AGCTATTACT	TTTACCCCTTC	CCATATCAC	TAAGTTCCA	CCTAGAGCCT	TGAATGGTT	6000
ATCAGTCGTC	TTTCGGAAAT	GTTCGGGATC	TGGTACAAAT	TTTCAATCA	CATCTGTATA	6060
TTTTTAATT	TCCTCATCAG	AGAAAGTATT	TTTCATCAGA	GACAAGGTAT	TGTTGATGCC	6120
ACGAGGTGTA	CCGATTTCAT	AATCCCACCA	GTTCACCAACA	ATGCTTTTT	CACTATTGTA	6180
GACATGTTA	TGCATCCATT	CCATGGAATC	CCTGACTGTT	CGAACGACAG	TTTCATCTTG	6240
ATAATAACGA	GAAGAAGGAT	TGGTCACTTG	CTTGGCCATC	TCCTCCAATT	TCCGATAAGT	6300
GGCAGTCAGA	TTTGCAGACG	TTTTATAATT	TGAAAATTTT	TCCCACAAAT	AGGTGCGGTC	6360
CGCCTGACTT	GAAAATCTGG	ATAGGCTATC	AGCTACCTTT	CCTTCCAATT	CCTGGTTAA	6420

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TTTGGCCATC	TGTTCATTTT	TAGAACATA	GTATTGATTG	CCAGCGATGA	TGCCATTCCA	6480
GTCATCCAAA	CGGTCTGTGT	ATGCATCCTT	AACAGAGGCC	AGAATCTCA	AAGGAATCTT	6540
TTTCACCTCC	TTGCCATCTT	TACTGACAAT	GACATTGGTT	GTCCCTTCCT	TAAGAGGTT	6600
TAAAATCCA	TTTTGACTG	AAGAACCGTC	AGGATTTCT	ACCTTATAAG	TATAGTCCGC	6660
AAGAGAAAAA	ACATGTTTT	TTCCAATTGG	AAATCAATC	TTTCCTCAA	GCTGTTTATC	6720
TGTTTGAGAA	TCCTCAGAAA	GCTGGTCTGC	TACCTCTACC	AGCTCAATAT	CCTTAAAGGA	6780
AACAGTCCCA	GTTCTGTGTT	CATAGAATAA	CTCCAGCTTG	ATTTTATCAA	CATCTAAAGT	6840
CGGGCTATAG	TCTGCTCAA	TGGTCTGCCA	GTCCTTGTG	CCTGACGTCG	TTGCAGAATT	6900
CCACAAATCGC	TTGTCCCTAC	CACTTCCTC	AATGATACGA	ACTTTGGCAA	TCCCAGTTTT	6960
ATTATCTGTT	TTAACCTTGA	AACGCAGTTT	ATACCTTTTC	TTAGCTCAA	TAGGAACCAT	7020
ACGGTGAAGC	GCTGCCCTTA	ATTTCTCATG	GCTTGAGATA	GTGATAGCCC	CATCCTTAGC	7080
CTCAATGACT	CGAGTTGAGG	CATCTGCACT	ATTCTCTGG	TCTACCCAAG	CTGACCACCC	7140
CCTGAGCTTT	GCTTCCGTGTC	CGG				7163

(2) INFORMATION FOR SEQ ID NO: 68:

- (i) SEQUENCE CHARACTERISTICS:
- (A) LENGTH: 9244 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 68:

CGTTATAACA	TACATGTAAG	CGGTACCCAA	AATGGTGCCA	AGTCAAAATT	TTTAAGGAGG	60
AAAATACATG	TCCTCACATC	CAATTCAAGGT	CTTCTCAGAA	ATTGGGAAAC	TGAAAAAAAGT	120
TATGTTGCAC	CGTCCAGGCA	AGGAGTTAGA	AAACTTGTG	CCGGACTATC	TTGAAAGGCT	180
TCTTTTGAT	GATATTCCCTT	TCTTGGAAAGA	TGCTAAAAAA	GAACATGATG	CATTTGCCCA	240
AGCTCTCGC	GATGAAGGAA	TTGAGGTTCT	CTACCTAGAA	CAAATCGCTG	CTGAATCATT	300
GACCTCTCCA	GAAATCCCG	ATCAATTAT	CGAGGAATAC	TTAGACGAAG	CCAACATCCG	360
TGATCGTCAA	ACCAAGGTTG	CTATTCTGTA	ATTGCTTCAC	GGCATCAAGG	ACAACCAAGA	420
ATTGGTTGAA	AAAACAATGG	CTGGGATTCA	AAAAGTTGAA	TTGCCAGAAA	TTCCCTGACGA	480
AGCTAAAGAT	CTAACTGACT	TAGTTGAATC	AGAGTATCCA	TTTGCATTG	ACCCGATGCC	540
AAACCTCTAT	TTCACTCGCG	ACCCATTGC	AACAATTGGA	AACGCCGTAT	CGCTTAACCA	600
CATGTTGCA	GACACTCGTA	ACCGTGAAAC	ACTCTACGGT	AAGTATATCT	TCAAATACCA	660

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CCCAATCTAT GGCGGAAAAG TGGATTTGGT CTACAACCGT GAAGAAGATA CGCGTATCGA	720
AGGTGGAGAC GAGTTAGTTC TTTCTAAAGA CGTCCTTGCA GTAGGTATCT CTCAACGTAC	780
AGACGCAGCT TCTATCGAAA AACTTTGCGT CAACATCTTC AAGAAAAATG TTGGCTTCAA	840
GAAAGTTTG GCCTTGAAAT TTGCTAACAA CCGTAAATTG ATGCACTTGG ATACTGTCTT	900
CACTATGGTA GACTATGACA AGTTCACTAT TCACCCAGAA ATCGAAGGCG ACCTTCACGT	960
TTACTCAGTT ACTTACGAAA ACGAAAAACT TAAAATCGTT GAAGAGAAAG GTGACTTAGC	1020
TGAACCTCTT GCTAAAACC TTGGTGTAGA AAAAGTTCAT TTGATTGTTT GCGGTGGTGG	1080
CAATATCGTA GCAGCTGCC GTGACAATG GAACGACGGT TCTAACACTT TGACCATCGC	1140
ACCTGGTGTG GTAGTTGTTT ATGACCGCAA TACCGTGACC AATAAGATTT TGGAGAATA	1200
CGGGCTTCGC TTGATTAAGA TTCCGGAAAG TGAATTGGTT CGGGGCCGTG GTGGACCTCG	1260
TTGTATGCT ATGCCATTG AACGTGAAGA AGTGTAAATCG CTGTTGATA TTGTCATAATA	1320
GAAAATGTA AAAATAGAAA GAGGAATAA TAAAATGACA AATTCACTAT TCCAAGGACG	1380
CAGCTCTTA GCAGAAAAAG ACTTTACCCG TGCAAGGTTA GAATACCTTA TTGGTCTTTC	1440
AGCTCACTTG AAAGATTTGA AAAACGCAA TATTCAACAC CACTACCTTGC TGGAAGAA	1500
TATCGCTCTC CTATTGAAA AAACATCTAC TCGTACTCGT GCAGCCTTA CAACTGCGGC	1560
TATCGACCTT GGTGCTCACCC CAGAATAACCT CGGAGCAAAT GATATTCACTG TGGGAAAAAA	1620
AGAATCTACT GAAGATACTG CTAAAGTATT GGAGCTATG TTTGACGGGA TTGAATTCCG	1680
CGGATTCAAGC CAACGTATGG TTGAAGAATT GGAGAATTC TCAGGCCTTC CAGTATGGAA	1740
CGGTCTAACT GACGAATGGC ACCCAACTCA AATGCTCGCT GACTACTTGA CTGTTCAAGA	1800
AAACTTCGGT CGCTTGGAAAG GCTTGACATT GGTATACTGT GGTGATGGAC GTAACAACGT	1860
TGCCAACAGC TTGCTCGTAA CAGGTGCTAT CCTTGGTGTC AATGTTACA TCTTCTCAC	1920
AAAAGAACTC TTCCCAGAAA AAGAAATCGT TGAATTGGCA GAAGGATTTG CTAAAGAAAG	1980
TGGCGCACAT GTTCTCATCA CTGAAGATGC TGATGAAGCA GTTAAAGATG CAGACGTTCT	2040
TTACACAGC GTTGGGTAT CAATGGGTGA AGAAAGACAA TTCGCAGAAC GTGTAGCTCT	2100
TCTTAAACCT TACCAAGTCA ATATGGACTT AGTTAAAAAA GCAGGCAATG AAAACTTGAT	2160
CTTCCTACAC TGCTTGCCAG CATTCCACGA TACTCACACT GTTTATGGTA AAGACGTTGC	2220
TGAAAAAATTG GGTGTAGAAG AAATGGAAGT AACAGACGAA GTCTTCCGCA GCAAGTACGC	2280
TCGCCACTTC GATCAAGCAG AAAACCGTAT GCACACTATC AAAGCTGTTA TGGCTGCTAC	2340
ACTTGGTAAC CTTTATATTC CTAAAGTATA ATTGAGATA ATAAACCGTC TACCAACAGC	2400

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TATGAGGGCT GCGACTAATA GCTTAGTCC GGTCTCTTT TATGTAATGG TAATCTATTA	2460
TTTCTTATAA AATATGTCAA AAATCATTAA ATTGAAATCT AAACGCATTC TATTGAGTGT	2520
GATAAAGGAG AATTATGGC AAATCGAAA ATTGTAGTAG CTTGGGGAGG AAATGGGATT	2580
CTTTCTTCTG ACCCATCAGC AAAGGCTCAA CAAGAAGCTT TAGTGAAAC AGCTAAGCAT	2640
CTTGTAAAAAT TGATTAAGGGAGATGAT CTGATTATCA CTCACGGTAA TGGACCTCAA	2700
GTTGGGAATC TCTTGCTCCA ACATTTGGCA TCAGACTCTG AAAAGAACCC TGCCCTCCCA	2760
CTCGACTCAC TTGTCGCTAT GACAGAAGGT AGCATCGTT TCTGGTTGAA AAATGCTTTG	2820
CAAATGCTC TCTTGATGAA AGGCATCGAA AAAATGTTG CCTCTGTTGT AACGCAAGTT	2880
GTCGTAGATA AAAATGATCC AGCTTTGTT AACTTGAGTA AACCAATCGG TCCTTCTAT	2940
TCAGAAGAAG AAGCAAAGC AGAAGCCGAA AAAAGCGGAG CGACTTTCAA GGAAGATGCT	3000
GGCCGTGGCT GGCAGTAAGGT CGTTGCCCTCA CCAAAACCTG TTGACATCAA AGAAATTGAA	3060
ACCATCCGTA CTCTTTAAA TAATGGTCAA GTGCGCTGTAG CTGCAGGTGG TGGCGGTATT	3120
CCCGTCGTCA AAGAAAACAA TGGACATTTG ACTGGTGTGCG AAGCGGTTAT TGATAAAAGAC	3180
TTCGCTTCCC AACGTTGGC AGAATTGGTT GATGCAGACC TCTTCATCGT TTTGACAGGT	3240
GTAGATTATG TATTTGTTAA CTACAACAAG CCAAACCAAG AAAAATTGGA ACATGTGAAT	3300
GTTGCCAGC TGGAAGAATA TATCAAACAA GATCAGTTG CACCAAGGTAG CATGCTTCCA	3360
AAAGTAGAAG CAGCTATCGC TTTGTCAAT GGTGTCAG AAGGAAAGC AGTTATTACT	3420
TCCCTTGAAA ATCTAGGCGC CTTGATTGAA TCTGAAAGCG GAACAATTAT TGAAAAGGA	3480
TAAGTTGTT TACTAATAAG ATGTATTCTA TTTCTAGTAT CTTTATATCA AATTAGAAAAT	3540
TATTCTGAA AACATGTACA ATATTCAAA AGATACTAGT TTTAGACTTT AATATGGTAA	3600
AACAAATATA AATAGAAAGC GTTTCTTGA ATGTTTATTT AAGAAAGTAG TTGGTTTTT	3660
ACACTTTGTT AGACATCAGG AGGAAAACA AATGAGTGA AAAGCTAAA AAGGGTTAA	3720
GATGCCCTCA TCTTACACCG TATTATTGAT AATCATTGCT ATTATGGCAG TGCTAACTTG	3780
GTTTATCCCT GCGGGGGCCT TTATAGAAGG TATTTACGAG ACTCAGCCTC AAAATCCACA	3840
AGGGATTGAG GATGTCCTCA TGGCACCGAT TCGGGCTATG CTAGGTACTC ATCCAGAGGA	3900
AGGTTCGCTC ATTAAAGAAA CGAGCGCAGC GATTGATGTA GCCTTCTTCA TCCTTATGGT	3960
TGGTGGTTTC CTTGGCATTG TCAACAAAAC TGTTGCTCTT GACGTAGGGA TTGCTCTAT	4020
CGTGAAGAAG TATAAGGGCC GCGAAAAAAAT GTTAATTTG GTACTGATGC CTTTGGTTGC	4080
CCTCGGTGGT ACAACTTATG GTATGGGTGA AGAAACAATG GCCTTCTATC CACTCCTTGT	4140
GCCAGTTATG ATGGCCGTTG GTTTGATAG CCTGACTGGT GTTGCAATTAA TTTTGCTCGG	4200

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TTCTCAAATC GGCTGTTGG CATCTACTCT GAATCCATT GCGACAGGTA TTGCTTCAGC	4260
GACTGCCGGA GTTGGTACAG GGGACGGTAT CGTACTTCGT CTGATCTTCT GGGTTACCTT	4320
GACTGCTCTT AGTACTTGGT TTGTTTACCG TTATGCGGAT AAGATTCAAA AAGATCCGAC	4380
TAAGTCACTG GTTTATAGTA CTCGCAAAGA AGATTTGAAA CACTTTAACG TAGAAGAAC	4440
TTCATCTGTA GAATCTACAC TTAGCAGCAA ACAAAATCA GTTCTCTTCT TATTTGTGTT	4500
GACATTCATC TTGATGGTAT TGAGCTTCAT TCCATGGACA GACCTTGGCG TTACCATT	4560
TGATGACTTT AATACTTGGT TGACTGGTCT TCCAGTTATTG GTTAATATTG TCGGTTCATC	4620
TACTTCTGCA CTAGGTACTT GGTACTTCCC AGAAGGCAGCA ATGCTCTTTC CCTTTATGGG	4680
TATCCTGATT GGTGTTATTG ATGGTCTTAA AGAAGATAAG ATTATCTCTT CCTTCATGAA	4740
TGGTGCTGCT GACTTGCTCA GTGTTGCCTT GATCGTAGCG ATTGCTCGTG GTATTCAAGT	4800
TATCATGAAC GACGGTATGA TTACCGATAC AATCCTCAAC TGGGGTAAAG AAGGCTTGAG	4860
CGGTCTATCT TCACAAGTCT TTATCGTTGT AACTTATATC TTCTATCTAC CTATGTCATT	4920
CTTGATCCCC TCTTCATCTG GTCTGCCAG CGCAACTATG GGTATCATGG CTCCACTTGG	4980
AGAATTGTA AATGTCGTC CTAGCTTGAT TATCACTGCT TACCAATCTG CTTCAGGTGT	5040
CTTGAACCTG ATTPGCACCAA CATCTGGTAT TGTGATGGGA GCTCTTGAC TTGGACGTAT	5100
CAACATTGGT ACTTGGTGG AATTCAATGGG CAAACTCGTA GTCCCTATTA TTGTAGTGAC	5160
CATCGCCCTT CTTCTCCPTG GAACCTTCCT TCCATTCCCTA TAAAATAGTG AGTGAGGTGA	5220
TTCCATGAAA ATAGATATAA CAAATCAAGT TAAAGATGAA TTTCTTATAT CATTAAAAAC	5280
CTTGATTTCC TATCCTTCAG TACTCAATGA AGGAGAAAAT GGAACACCTT TTGGACAAGC	5340
AATCCAAGAT GTCCCTAGAAA AAACCTTTAGA GATTTGTGCA GACATAGGTT TCACTACCTA	5400
TCTTGACCCCT AAAGGTTATT ACGGATATGC AGAAATCGGT CAGGGAGCAG AGCTTCTGGC	5460
CATTCTCTGT CATTGGATG TTGTTCCATC AGGTGATGAA GCAGATTGGC AGACACCGCC	5520
ATTTGAAGCA ACTATCAAAG ACGGCTGGGT ATTGGACGT GGTGTCAG ATGATAAAGG	5580
CCCTTCGCTC GCAGCTCTCT ATGCAGTAAA AAGCTTGCTG GACCAAGGTA TTCAGTTCAA	5640
AAAGCGCGTA CGCTTATCT TTGGTACCGA TGAGGAAACC CTCTGGCGCT GCATGGCACC	5700
CTACAATACC ATCGAAGAAC AGGCCAGTAT GGGCTTGCA CCTGACTCAT CTTTTCTCT	5760
GACCTATGCT GAAAAAGGCC TTCTACAGGT CAAACTTCAT GGCCTGGAT CGGATCAACT	5820
AGAGCTTGAA GTAGGAGGCG CCTTTAACGT TGACCAGAC AAGGCCAACT ACCAAGGTCT	5880
CCTCTATGAA CAGGTTGTA ACGGTCTCAA AGAAGCTGGT TATGATTACC AAACCACTGA	5940

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ACAAACCGTA ACGGTTCTCG GAGTGCCAAA GCATGCTAAG GATGCTAGTC AAGGTATCAA	6000
TGCTGTCATC CGACTAGCTA CCATTCTTGC TCCTCTCCAA GAACACCCCTG CTCTCAGTTT	6060
TCTTGCACA CAAGCAGGTC AAGACGGCAC AGGAAGACAA ATCTTTGGTG ATATAGCAGA	6120
TGAAACCTTCT GGTACACCTAT CCTTTAATGT CGCAGGTCTC ATGATCAATC ATGAACGTTC	6180
TGAAATCCGT ATTGACATTC GGACTCCTGT CTTAGCTGAC AAGGAAGAAC TAGTAGAGTT	6240
GCTTACAAAGA TGTGCACAAA ACTACCAACT CCGCTACGAA GAGTTTGACT ATCTAGGCC	6300
TCTATACGTC GCAGAAGACA GTAAACTCGT TAGCACACTG ATGCAAATCT ACCAAGAAAA	6360
GAATGGCGAT AACAGTCCTG CTATTTCATC CGGTGGTGCC ACTTTTGCTC GCACCATGCC	6420
AAATTGTGTA GCCTTCGGCG CCTTATTCCC AGGAGCGAAG CAGACAGAAC ATCAGGCAA	6480
TGAATGTGCC GTTCTAGAAC ATTTGTACCG TGCTATGGAT ATTTATGCCG AAGCCGTCTA	6540
TCGACTTGCA ACTTAATCAG GCAACTGTTT CTACCAAAAA AAATCGACCG ATTAATGAAC	6600
TGCACCCCAA AAGTTAGACA GAATAAATCT AACTTTGGG GTGTTTTATT ATGAAATTGA	6660
GTTATGAAGA TAAAGTTCAAG ATCTATGAAC TAAGAAAGCA AGGACAAAGC TTCAACAGC	6720
TTTCAAAAAG ATTTGGTGTG GATGTTCTG GTCTAAAGTC ATCTGAATCT TTGAGATGAG	6780
CTTTATAAAT CGCTTTTTTC AGTTTTTGCA CTGGTGTTC GATAAACTCA AACTTTTAG	6840
CCGTGGTATT GCCTGATTTT ATAGTATATT GAAACTAGAA TAGTACACCT CTCCTTCTAA	6900
AACATTTTA GAAATCGATT TGACTGTCCT GATCGATTG TCCTGTTCTT ATTTCAATT	6960
ACTATATTTG AGCCACTTCG TCTTTAACGG CTTTATTCTA AAGCTCTTGT AATTTTTCTT	7020
TAATCAAT TACTCTGAT TTTCCGTTGT AATTTATTGT AATAGGTTTT AACTTACCTA	7080
ATTTCTCGAC ACGCTCATTA ATTTGATCTT TTTTGAAAGGC TGCTTATGTT TTTCTAAGA	7140
TTTTTTCAAA AATATATTAA TCAGATAGCG GTTGTCTTC TTCTTCAGCT TGGTTTTGT	7200
ATTAATTGAA AACATAAGGA ACAAACTCTT CATAGTAACC TAATGCTCCC ATAAGTTCAA	7260
AAGCTTGTCTT TCTAATTCAA ACCATTGCAA CTCAGATTTG AGCTTTTCAG ATAAATCCTG	7320
CTCATCCAAA TAATGACTTG AAAATTAGTGC TGAACACTGTT TCTGTATCCT GTACAGGCTG	7380
AGCACCCATA CCAGCAAAAA ATAAACTCGT TCCTAGCAAG ACCGAACAAAG CTCCTATTGC	7440
ATATGGCTC AAAGAAAAAC GCTGCTTCT CTCAAATTGA AATTCTTCA TCCCCTCTCC	7500
CATCATTCAT TATTACTGTA TATTTTGAT ATCAGAAATA GTTGTATTC ACAAAATCTT	7560
CTAGTTATTC CCTTATCATT CCTAATTAAG GGAGATAACA TACAATAATT TTTAGTTAAA	7620
TGTATATCGA TGTTTTTGT TTTCTTAAT AAACGCAATA CAAAAGAGC CTGTTACCAA	7680
GCTCTTGTA CTCAATGAAA ATCAAAGAGC AAATTAGGAA ACTAGCCACA GGTTGCTCAA	7740

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AACACCGTT TGAGGTTGCA GATAGAACTG ACGAAgTCAG CTCAAAACAC TGTGGAGG	7800
TTGCAGATAG AACTGACGAA GTCACTAACCA TCTATACGGC AAGGCGACGC TGACGTGGTT	7860
TGAAGAGATT TTCGAAGAGT ATTAGTCTAT TATTTCTCT CAGCGCGAAG GGCTGACAAG	7920
ATTTGTGTTG GGATATCATC CACACCATTG GGAGTATTTG GTAAAAAGAT AGTTTGATT	7980
CCTTTAGAGG CAAAGGTATT CAAGGTATCC AAATACTGGT TGGTCAAGAG GATAGACATG	8040
ATTTGTTCTT CTGTCATGCC AACATTGGCT TCCMTGAGTT CGGTGATAGA CTCTGCCAAT	8100
CCATCCACAA TCGCCCTACG TTGTTGGCA ATCCCCACAC CATGAAGGCG GTCTTTTCT	8160
GCTTCTGCTT CAGCTGCAGT GACAATTAA ATCTTGTCAAG CTTCCGCCAA TTCTTGTGCT	8220
GCGACCCGCT TACGTTGCGC CGCATTGATT TCATTCATGG ATTGCTTAAC TTCTGCATCT	8280
GGTTCGACCT TGGTAATCAA GGTTTCACG ATAATGTAGC CGTAAGTGGT CATTTCCTCT	8340
GCTACTGGT GTTGAACCTC AAGGGCAATC TCATCTTTT TCTCAAACAA TTCATCCAAG	8400
GTAAATTGGT GAACAGAAGA GCGAAGAGCA TCTTCGATAT AAGATTTAAT CTGAGATTCT	8460
GGACGTATGA GTTTATAGTA AGCATCTGTC ACGCTCTGCT CGTTGACACG GTACTGAGTC	8520
GCTACATTCA TCATAACGAA CACATTGTCC TTGGTCTTAG TCTCAACCAC AATATCACTT	8580
TGCAACAAGC GCAACTGAAT CCGTGCTGCA ATCGAGTCAA TCCCAAAAGG CAAGCGAATA	8640
TGAATACCGC TATTAGCAAC CTTTTGGTAT TTCCCAAAGC GTTCAATAAT CGCCACCGAC	8700
TGCTGACGAA CCACATAAAC TGTACTCAGT GTGACTATCA CCAATAGGAGCACACAAACA	8760
ATCAGAAAAA TCATCAAAAAT TATTGCCATA ATGGAACCTC CACAAGTATT TTTCTAGTAT	8820
TATAGCACAT TTAAAGAAGG CTGTGCCGTT TTACTGCGA TTTTCTCTGA AATGTCAATA	8880
ATTAGAGGTG AATTGTCCTA TTGTCGTCCA ATCTCTTGCT AAAATAACTC TTTATAAAAG	8940
GCAATCGTTT CTTCTAAGGT TGGCATAAAAT GGATTTCTG GTGCGCAGGC ATCAATCAAG	9000
GCATTCTTAG AAAGGTATTC AAAGTCGAAA TCTTTTCTT CAATACCAAG TTCAGTCAGT	9060
TTCTTAGGAA TACCTACTGT CTCAGAAAGC TTCTCAATCT CAGCAATCGC ATAATCGGCA	9120
CATTCTGAT CTGATTTACC TTCTACATGA AGTCCCAGG CTTTGGCAAC ATTGCGGAAA	9180
GCTTCTGGTA CACGTTAGC ATTTTCACGT TCTATAACTG GTAGCAACAT GGCACAGCAC	9240
ACGG	9244

(2) INFORMATION FOR SEQ ID NO: 69:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 8898 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double

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(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 69:

GATCTGAAC	TTATCATCAT	AACTTAATT	CATAATAAA	ACACCCAAA	AGTTAGATT	60
TTTCTGTCTA	ACTTTGGGG	TGTAGTCAG	TCATTGGACT	GACGTTTTT	TGTATGCTTA	120
TTTGATTTG	ATGTAGTTGA	TACCATCTGC	TTTGGTGC	ACTGCTTTTC	CAAAGAAGGC	180
TGCTAACACA	AGAATTGTCA	AAACATAAGG	TGCAATTG	AGATAAACCG	CTGGCACTCC	240
TTGTAGGAAC	GGCAATTGAG	AACCATAAC	AGCCAAACTT	TGTGAAAGTC	CAAAGAAGAG	300
ACTAGAAAGC	ATAGCACCGA	TTGGATTCCA	TTTCCAAAG	ATCATCGCAG	CAAGGGCGAT	360
AAATCCAGGT	CCAACAAATAG	TTGTCACTGA	GAAGTAACT	GAGATTGATT	GCGCATAAAAT	420
CGCTCCGCCA	ATTCACCTA	AAAAACCTGA	ATAATAACC	CCTAAATATC	TCATCTTGTA	480
GACGTTGATT	CCCAAGGTAT	CCGCTGCTTG	AGGATGTTCA	CCGACAGAGC	GGAGACGAAG	540
ACCAAATTGA	GTCTTAAAGA	GAATAAACCA	AGCAAGGAAT	GAGAAGGCAA	TCGCCAGATA	600
ACCAAGTAGA	CTAGTTGACT	TGAAGAAGAT	ATCACCAATC	ACTGGGATAT	TTGCCAAGAC	660
TGGGAAATCA	AAGCCTCCAA	AAGTTGACT	TAGGTTGTCG	GTGTCCTT	TGTTATAAAAG	720
AACTTTAACT	AAGAAAACAG	CCAAGGCAGG	CGCCATCAAG	TTCAATACCG	TACCGCTGAC	780
AACATGGTCT	GCACCGAAAT	GAACCGTCGC	TGCTGCGTGG	ATGATAGAGA	AAACACTTAC	840
AACCAATCCT	GCTACAAGCA	AGGATAGCCA	TGGAGTTGCT	GCTCCAAATT	GTTCTGCAA	900
TTCAAGGTAA	AAGACAACCTC	CAGAAAAGGC	ACCCATAACC	ATAATTCCCTT	CAAGGCCAAC	960
GTGTTACCA	CCACCACTGTT	CAGAGAAAAC	ACCACCGATA	CTTGTAAGA	TGAGAGGTGC	1020
TGAGTAAATC	AGCATAGAAG	ACACCAAGAG	GGGGAGCAAG	GTTATAATAG	ACATCTTAC	1080
TTACCTCCCTT	TAACCTGTTT	TTTCGGTTTG	ACAAAGCGTT	CGATAAGGTA	ATGAACACTG	1140
ACAAAGAAGA	TAATAGACGC	TGTTACAATG	CTGACAAGCT	CAGATGGTAC	CTGCGCCGCA	1200
TTCATACCAG	GAGCCCCAAC	TTGGAGAACG	CCAAATAGGA	AGGCTGCAA	GAGTATACCA	1260
ATTGGTGAAGT	TGGCCGCAAG	CAAACAAACC	GCCATTCCGT	TAATCCGAT	AGCTAATGAC	1320
GAACCTTGAA	CATAGACGTT	CTGGAAGGTT	CCCAAACCTT	CAACAGCTCC	ACCAAGACCT	1380
GCCAAAGGCAC	CTGAAATAAT	CATAGATAGG	ATAATAGTCC	GCTTGGCAGA	AAATACCAGCA	1440
TATTCCTGAAG	CATGTGGATT	AAGACCAACT	GCACGGATT	CAAACCAAG	AGTTGTTTC	1500
TTGAGCATGA	ACCAAATAAC	TGCAACGGCA	ATGATGGCA	AGAAAATACC	AAATATTCA	1560
CGTGAGTTAC	CAGTCACCTC	AGCCAACCAA	GGTGTCTGAT	AGGTTGCATT	AGCCCCAAC	1620

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CGAATGGTCG AATCTGTACT TTGCATGAAG TCTTTAGGGA AAGCATGGAT AAAGGCATTC	1680
CCTACATACA AGACAATGTA GTTCATCATG ATGGTTACAA TAACCTCTGA CGTCCCTAGA	1740
TAGGCCCTAA GAATACTCGG AATCGCTCCG ACAATCCCAC CAGCAATCAA GGCAATCACG	1800
ATGGTTGCTA GAATCATCAA GGGACGGGGC ATATCTGGAT GCGACAGGGC AAACCAACCA	1860
CTGAGAACATCC AACCTGCCAA AGCCTGACCA GGAAGTCCGA CGTTAAAGAA ACCAGCTCGA	1920
CTGGCAACGG CAAAACCAAG ACCAACATCAAG ACCAGAGGAC CCATAGCACG GAAGATTCT	1980
CCAATCCCAC GCAGACTGCC AAAGGCTGTA TAGAACATT CTTCTGTAGCC CCAAATAGCA	2040
TCATAACCGA AGATCCACAT GACAATGGCT CCGAGTAAAA TTCTCTAGAA TACAGAAATC	2100
AAGGGAACCG AAATTTGTTG TAATTTTTTA GACATCACTC TTCTCCTTTC CCAAGTTCC	2160
ACCAGCCATC AAGACACCAA GTTCTTGTTT ATPTGGTTGTT TCTGGTGATA CAATACCTTG	2220
AATCTTACCA TCGTGGATAA CGGCAATACG CTCTGAGACG TTTAAAATCT CATCCAATT	2280
AAAGCTGACA ACAAGGACAG CCTTGCCATT ATCACGCTCT TCAATCAAGC GTTGTGGAT	2340
ATACTCAATG GCACCGACAT CCAACCCACG AGTTGGCTGG CTAACGATAA GGAGATCAGG	2400
ATCTCGATCA ATTTCACGAG CAATAATTGC TTTTTGTTGA TTTCCTCCTG AGAGTGCAGC	2460
TGCAGGAACCT AATTCACTGG CAGCGCGAAC ATCAAACCTCT TCCATCAGCT TTTTAGCATA	2520
AGAAGTAATA TTTGAATAAT TCAAAATTCC ATTTTTACTA TGTGGTTCTT TATAGTAGGT	2580
TTGAAGGGCA ATATTTTCAG ATATCATCAT TTCCAAAATC AAGCCATCAC GGTGACGGTC	2640
TTCTGGAACG TGCCCAACAC TTAGTTCTGT AATCTGACGT GGGTGCAAGC CTACAATTGA	2700
ATCTCCTTT AGCTCAATGC TACCAAGATTC AACCTTACGA AGACCTGTAA TGGCTTGAAT	2760
CAGTTAGAC TGACCATTTTC CATCAATCCC CGCAATACCA ACAATCTCTC CAGCACGAAC	2820
ATCCAAGGAC AGATTTTTAA CAGCTGGAAC ACCACGGTTT TCATTGACCA CCAAATCTTT	2880
GATAGACAAA ACCACTCTT TTGGTTTAGA GGCTTGCTTC TCTGTTTAA AGGAAACAGA	2940
ACGCTCTACC ATCATTTCCG CCAAATCAGC ATTGGTAGCC CCTGCAATT TCAACGGTTTC	3000
AATTGATTTTC CCACGACGGA TAACTGTAAC ACGGTCAGAA ACTGCTCGAA TTTCATCCAA	3060
TTTGTGGTA ATCAAGATAA TTGATTTCC TTCTTGACCA AGATTTTTCA TAATAGCCAT	3120
CAACTCATCA ATTTCTGATG GAGTCAAAC AGCCGTTGGT TCGTCAAAGA TAAGGATATC	3180
AGCCCCCGA TAAAGTGGTT TTAAAATTTC TACACGTTGT TGGGCTCCAA CTGAGATATC	3240
TGCTACCTTG GCAGAAGGGT CAACAGCTAA GCCATAACGT TCAGAAAGAG CCTTGATTTC	3300
TTTGCTAGCT CCAGCGATAT CTAGCACACC ATTTTAGTC AATTCACTAC CTAAAATGAT	3360

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GTTTCAGCC ACTGTGAAGG CTTCAACCAA CATAAAAGTGC TGGTGAACCA TCCC GATTCC	3420
CAAGCTAGCT GCTTTAGATG GGGAGTCGAG ATTGACA ACT TGACCGTTGA CGCG GATTTC	3480
ACCACTAGTT GGTCAGAAGA GGCGCTGCTAA CATGTTCACT AGCGTGGACT TACCAGCCCC	3540
ATTTTCCTCT AAAAGTGCAT GAATTCACC TTTTCGTAGG TGCAAGTTGA TTTTGTCTGTT	3600
GGCAACAAAT CCACCAAAACA CCTTGGTAAT ATCACGCATC TCAATGACAT TTTCGTGTGC	3660
CATGTGCTCT TCCTTCAGA GTCTTATTTT ATTTCAATAA AACTTGCTAG TTTGTCTAGT	3720
AGCAAGCTTT ACTTAGACAA AATGACTTTG TCTCAACTCT TAAAAAAAGCG GCCCTTGGCC	3780
GCTTCCTAAG AAATGACTTC CATCCATTAT TTTTCAGGAA CTTTTACGCT TCCATCAAGG	3840
ATTTTAGCTT TTGCACTTC GACAGCTTTT TTACCTTCTT CTGAAAGGTT TGTTACTGCC	3900
AAGTCAACCC CTTTATCCTT CAATGAGTAA ACGATCACTT GACCGCCAGG GAATTCTCCT	3960
CTTTCTGCCT TGTTAGAAAAT ATCTTTACA GTTGTACCA CTTGTTCAA AGTAGATACA	4020
AGAACAAAGT TTGATTCTTT GCCATCTTTA GAAGTGTATT TACCTTCTGC TTCTTGGTCA	4080
CGATCAACAC CGATAACCCA AACTTTTCA TTTTCAGGAC GGCTTTCGTT GAGAGATTTT	4140
GCCTCTGCAA AGACACCTGC ACCTGTACCA CCAGCTACTT GTAAACAAAT ATCTGCACCG	4200
GCTGCGTATT GTGCGGCTGC AATTGTTTTA CCTTTAGCCG CATCACCAAA TGAACCAGCG	4260
TAGTCAACTT GGACTTTGAT AGATGGGTCT ACTGACGCAA CACCAAGCTT GAATCCTGCT	4320
TCAAAACGAG AGATAACTTC AGATTCGATA CCACCTACAA AACCAACTTG TTTTGTCTTA	4380
GTTGTTTTG CTGCAGCCAC ACCTGCAAGg TAACCTGACT CATTATCAGC GAAAGTTACG	4440
CTCGCAACAT TCTTTGGTC TTTAATCACA TCATCAATCA AGACATAGTT CAAGTCAGTG	4500
TGTTCTTTG CTGCATCTTT AACTGCATTA TTAAGGGCAA AACCAACACC GAAGATTAGG	4560
TTGTAACCTTC CAGCCGCTTG TTGCAAGTTG TTAGCGTAGT CAGCTTCACT TGTTGATTGG	4620
AAGTAAGTGA AACCGTTATC TTTTGAAAGA TTGTTGTTCTT TACCCCAAGC CTGCAAACCT	4680
TCCCAAGCTG ATTGGTTGAA TGATTTGTCA TCAACACCAC CAGTATCAGT GACGATTGCT	4740
GCTTTGTCT TCACATCAGA AGATGAAGCT GCGTTACGAG AAGAGCGGTT ACCACATGCA	4800
GCAAGTCCAA CTGCTGCCAC TGCAACTAGG CCAAGACCTA GCCATTGTTT CTTGTTCACTT	4860
ACTGAACCTC CTAATAAGA TGTGCAACGA TGTGCAAGT ATGGATTGGT TGGCCACAAG	4920
GACCGTGCCA CTCAGAGAGC GACTCAGACT AGTTAACGTC TGTAAAAGAG TATGGAAGTA	4980
ATTCGGGAC CGTCATCTCG ACCGTCGATT TATCTTTGC GACTAAGGTC ACTTTTAGAT	5040
CTTGTTCAAA AAATTCAAGCC ATCACTTGGC GACAAGCACC ACATGGCGAG ATCGGTTTT	5100
CAGTTTGACC ATAGACAATC AATTCTGAAA ATTCTCTTTG GCCTTCAGAT ATAGCCTTAA	5160

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AAATAGCTGT TCTCTCACCG CAATTGGTCA AAGGATAGCT AGCATTTCAT	5220
CCGTGTAAC ACTTCCGTCT TTAGCTACTA AACTGCTCC GATAGGAAG TGAGAATAGG	5280
GGACATAGGC ATGTTGCTG GTTCAATTG CCAGTCAAT CAACTCAGTA GTCGCCATCT	5340
GCCAATTCTC CTTTAAAAT AGCTACCCCA GCTGACGTTG CGATACGGGT CGCACCTGCT	5400
TCGACAAAGG CAAGAGCATC TGCTATAAGAA CGAGCTCCAC CGGGCGCCCT GACACCCATA	5460
TCAGATCCAA CTGTTTCACG CATTAAATGTA ACATCTGCTA TCGTAGCACC ACCAGTTGAA	5520
AAGCCAGTAG ATGTTTGAC AAAGTCAGCC CCAGCTTTT GGGCCAATTG GCAAACAACA	5580
ACTTTTCTT GGTCTGTCAG AAGGCAAGCT TCAATAATGA CTTTCACTAA CTTATCACCA	5640
CTTGCTTCCA CTACTGCGCG AATATCTGAC TCAACCAAGG CTAAATTACC TGATTTGAGA	5700
GCTCCAACAT TGATCACCAT ATCAATCTCA TCTGCACCAT TTTGGATAGC TTCTTTGTC	5760
TCAAATGCTT TCACGGCTGA AGTGTGGCT CCCAAAGGGA AACCTACTAC TGTGCAAACC	5820
TTAACATCTG TGCCCTCAAG TCCCTTTTTA GCATGTTCAA CCCAGGTCGG ATTAACGCAA	5880
ACACTGGCAA AGTCATACTC TCTAGCCTCA GACAACAAAC TATCAATTG TTTTTTCTTT	5940
GCATCTTGTG TTAAAGCGT ATGATCTATA TATTTATTAA ATTTCATTTTC GGTTTCCCT	6000
CCATTTAGGA GATGATTCT ACAATTTCAC GGATTTTTT CACTTCATCA CTTATTTTAA	6060
CACATTTTG GAAATCTGTA ACTAGTTGAG GTGGAATTTC TTCATTTGT TATACTTTG	6120
CAACAAATTTC ACCCTTTGAG ACGGAGTCTC CAATCTTCTT TTCAAAAACA ATTCTGTGTT	6180
CATAGTCCAA GGCAATCAGAC TTAACTGCAC GACCAGCACC CAGCCTCATG GCATAAAGAC	6240
CAAAGTCCAT AGCTGGAAGA CCTGAAATGA CACCCGTTTC CTGAGCAGGG ATTTCCACCA	6300
CATGAGCTAC ATTTACAGGA CGATAGAGGT CTTCCAAGTC TCCACCTGG GCTTGCACCA	6360
TTTCCTCAAA CTTAGCCAGT GCTTGACCAT TCTCAAGATG TTGGTGAACCT TCTTCAACAG	6420
TTTTGTTAAC ATTTGCCAAA CCAAGCATAA TTTGAGCCAA TTCACAAATA AAGTGGTAA	6480
TATCCTGACG TCCTTGACCT TGCCTAAATCT CCAATGCTTC AAGGATTTCC AGACGATTTC	6540
CAATCGCTCG TCCCAAAGGC TGGCTCATAT CCCTAAATCAC TGCCTACTGTC TTCCGTCCAA	6600
CAACCTTACCA AAGATCTACC ATAGTTTGAG CCAACTCAGC CGCCTCATCA ACCGTCTTCA	6660
TGAAGGCACC CTCACCGACA GTCACGTCTA GCAAAATAGC ATCCGCCCT GCCGCAATT	6720
TCTTGCTCAT CACCGAACTC GCAATCAAAG GAATCGTGT GACAGTTGCG GTCACATCAC	6780
GAAGGGCATA GAGAAGCTTA TCTGCTTTGA CCAGCTGGTC TGATTGCCCA ATGACAGATA	6840
CTCCAATATC CTGAACCTGA CGAATAAAAT CCTCTTGACT ACGTTCTACT TGATAGCCCT	6900

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TAATGGACTC	CAATTATCA	ATTGTTCCGC	CTGTATGCC	AAGACCACGA	CCACTCATTT	6960
TTGCTACAGG	CACACCGAAG	CTAGCAACAA	GAGGAGCTAA	AATCAAGGTT	ACCTTATCGC	7020
CGACACCACC	AGTAGAACATG	TTGTCACATT	TCACACCATC	AATGGCTGAC	AGGTCAAAC	7080
CTTGCCCCAGT	CTTAACCATA	TTCATCGTTA	AATCAGAGAT	TTCTCGAGTC	GTCATTCCCT	7140
TAAAATAAAC	AGCCATAGCA	AAGGCAGACA	TCTGATAATC	AGGAACAGTT	CCTGATAACAT	7200
AGCCTTCTAT	CAGCCATTCA	ATTTCACTTG	AAAGTCAGTTC	TTGACCGTCT	CGTTTTTTT	7260
GGATTTAACATC	AACTGCTCTC	ATTCTTTCAC	ACTTCTAAGG	ATATAGTATC	CCTTGTCTTT	7320
TTTAAGGATT	TCACAATTGC	CAAACACATC	TTCCATCTTA	GACTTGGCAC	TTGGAGCTCC	7380
TTGTTTTTC	TGGATGACGA	TGGTCAAATC	TCCACCAATT	TCCAAGAAAT	CTTTACTTTT	7440
CTCGATGATT	TCATGACGA	CTTGCTTGCC	CGCACGGATA	GGAGGATTGG	AAATGACATG	7500
GTCAAATCGC	CCTTGAACTC	TTGCATAAAAT	ATTAGATTGA	AATATCGTCG	CTTTGCATT	7560
ATTTTTTCA	GCATTCTCT	GAGCTAAATC	CAGGGCACGA	GTGTTAATAT	CAACCATGGT	7620
CGCCTGAAC	CCGTAACCT	TGACCAAGGA	CAAACCTAAT	GGACCATAAC	CACAGCCTAC	7680
ATCTAGGACT	GTCTCTCTT	GGTGACATC	CAGACACTTG	AGCAAGAGTT	GACTTCCAAA	7740
GTCAACCATT	TTCTTGCTAA	AAACACCCGC	ATCTGTCAA	AAAGTCATTT	TTTCTCCAA	7800
CAAGTCCACT	CTCAACTCAT	GAATGTCGTG	AGCAGCGTCA	GGATTTCTG	CATAGTACAT	7860
TTTACTCATG	ACACTATTT	ACCATAATTT	GACTCAAATT	GTAAATCGTT	TACAAATTGA	7920
TAATAAAACG	AAAAGACCG	AAGAAAGCAA	GTCACGAAGC	CATTTCCTTC	AATCTCTTC	7980
AACACTTATA	AATAAAAC	CATTAGAAC	TATAATATC	ACAGTCCAGA	AAAAACAAA	8040
AAAGTTATCA	TCTATAATCA	GGCAGATTAT	TATTCTATT	GCTTAACCTT	AAAATACTTT	8100
ATTATCAACA	AAATTCTAA	CAAAATGTTT	AGATAAAAGC	CCAACTGATA	CGTTTATGTC	8160
AGGATTCCA	AACTTGTCCA	AAAGTCGTAC	AAATCTCTA	GTGACATGTG	GAAGAAATAA	8220
CCCTCTGTG	CAATCCGTAG	GACTAAAAG	CAATAACTAC	CCGCAGCAAT	CCATTTCGTC	8280
CATCGTTTT	TAGTAAGAAA	GCAATTAAGA	ACGAACAAAT	AAAGACAGCT	GTTACAATAG	8340
CATGTTCCAT	CAAAAAAGTA	AAACCGTAAT	AGGTTTCCAC	AAAGCATCTA	CCATTATCTG	8400
CATTGGTTCC	TTTTATAAAA	GGTAAAGCAA	AACCTAAAT	AAAACAGAGT	TCCAATATGT	8460
AACGTTTAA	GATTTCATA	GTACACCTCC	TATAAGTTGT	GAACCTAAAA	GCCCCCTTTA	8520
TAAGCTTATA	AATCAGTAGA	ATCTATCTCC	TATTTCATCA	ATAAATTGAT	CACTTATACT	8580
ATATACCATT	GACTTACAC	ATTCAAGAAA	CCGCTTATT	TTTTAGCTT	TTTATGGTAT	8640
GATAGACAAA	ATATCTAGGG	GAAAACAAAT	GACCAACGAA	TTTTACATT	TTGAAAAAAAT	8700

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CAGCCGCCAG ACTTGGCAAT CTTTACATCG AAAGACAACA CCTCCTTG A	8760
ATTGGAATCT ATCAAGAGTT TTAATGACCA AATCAGTCTC CAAGACGT A	8820
TCTCCCCTTG GCTCATTTGA TTCAGATTAA CAAGCGA AAGGAAGATT TAGCCTTT	8880
AAAAGGAATT TTCCTCCA	8898

(2) INFORMATION FOR SEQ ID NO: 70:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 13188 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 70:

TATCTTAAACG aGGATGGGT TTATCGTCAG TCTTATTGCC CTAATTGTGG GAACAATCCC	60
TTAAATCATT TTGAAAAATAA TCGGCCTGTA GCAGATTTT ACTGTAATCA TTGTAGTGAG	120
GAGTTTGAAC TAAAGAGCAA AAAAGGAAT TTTTCATCAA CAATCAATGA TGGTGCTTAT	180
GCAACGATGA TGAAGCGTGT GCAGGCAGAT AATAATCCTA ATTTCTTTT TTTAACTTAC	240
ACAAAAAATT TTGAGGTAAA TAACTTCTT CTCCCTCCGA AGCAATTGT TACACCGAAA	300
TCGATTATTTC AAAGAAAACC ACTTGCACCA ACTGCTAGAC GAGCAGGTTG GATTGGTTGT	360
AACATTGATT TATCACAAGT ACCTTCTAAA GGAAGGATAT TTCTTGTGCA AGATGGACAA	420
GTTAGAGATC CAGAAAAAGT TACAAAAGAA TTTAAGCAAG GTTTATTTT AAGGAAGAGC	480
TCTCTGTAT CAAGAGGTTG GACAATAGAA ATTCTAAATT GTATAGATAA GATAGAGGGT	540
TCAGAATTAA CCCTTGAAGA TATGTATCGT TTTGAAAGTG ACCTAAAAAA TATCTTGT	600
AGAACAAATC ATATCAAAGA AAAGATTAGG CAACAGCTTC AAATATTAAG AGACAAAGAA	660
ATAATAGAAT TTAAAGGTAG AGGAAAGTAT CGGAAATTAT GAAAACGAAA CAACTGTG	720
CATCAGAAGA GGTGTATGAT TTCTTAAAG TCATCTGCC TGATTATGAA ACTGAAAGCC	780
GTTACGATAA CCTAAGTTA ATCGTCTGTA CCTTATCAGA TCCCGATTGT GTGAGATGGT	840
TATCTGAAAA TATGAAATTG GGTGACGAAA ACAACTAGC TTTGATGAAG GAAAAATATG	900
GGTGGGAAGT AGGAGATAAA TTGCCAGAGT GGCTACATAG CTCCTATCAT AGATTATTGT	960
TAATAGGTGA ATTATGGAA ACCAATCTAA AACTGAAAAA GTATACAGTA GAAATTACAG	1020
AAACTTTATC ACGTTTAGTA AGTATAGAGG CTGAAAATCC AGATGAAGCC GAACGACTTG	1080
TAAGAGAAAA GTATAAGAGT TGTGAAATTG TTCTTGTGATGC AGATGATTTT CAGGACTATG	1140

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ACACTAGCAT ATATGAATAG GTAGATGTTT TTATTTGTC AACAAAAAAG AGGCTCGCAC	1200
CTCTTTTCT TATTCTTTT TATGATTTAA TACGGCATTG AGGACAATAG CGAGTAGGCT	1260
GGCTACGACG ATTCCGTTG AGAAGAACAT TTGGAAGGCT GTCGGCATGC TGACAAAGAG	1320
ATTACTGTTG TTGAGACCGA CACCTGCAGC GATTGAAACA GCTGCATAA GGAAGTTGTG	1380
TTCATTGTTA GCAAAGTCAA CACGGGCGAG GATTGCATC CCTTGAAATG ATACAAAACC	1440
AAACATTACC AGCATGGCAC CACCGAGGAC GGAGCTTGGG ATGATTGGG CAAGGGCGCC	1500
AAACTTAGGA ACCAGTCCAA GGAGAACCGAG GAAACCAGCT GCGTAGTAGA TTGGCAGGCG	1560
TTTTTGATG CCTGACAATT TAACCAAACC AACGTTTGT GAAAATCCGG TGTAAGGGAA	1620
GGTGTAAAG ATTCCCTCGA GAAGTACGGC CAAACCTTCT GCGCGGTATC CGTTGCGAAG	1680
GCGCGTGTG TCGATTGGAT CCTTTGTGAT ATCAGACAAG GCCAGATAAA CACCAGTTGA	1740
CTCAACCATA GACACCCTTG CGATGATACA CATCATGACA ATAGATGAGA TTTCAAAGGT	1800
TGGCATCCC AAGTAGAGTG GAGTTGGAC ATGGACAAGT GGAGCTACCG CAACAGGAGA	1860
GAAGTCCACC AAGCCCATAG TAGCAGCAAT GGCAGTTCCA ACAACCAGAC CAATCAAAT	1920
AGAGATAGAC TTGATAAAATC CTTGGTAAAG GATGTTGATC AAGAGGATAA TCAGAACAGT	1980
AATAGCTGCA AGCAAGAGAC TTTGACCAAGT TGGCTCTGGG ACGTTATTTCC CCATATTTCC	2040
AATAGCGACA GGGATCAAGG TAAACCAAT CGTGGTAATA ACAGATCCTG TTACGATAGA	2100
TGGGAAGAGA TTGGCTACTT TTGAGAAGAT GCCTGAAACA AGAACACAGT AAATCCCAGA	2160
TGGCGATAAGG GCACCAAACA TAGCGCCACT ACCATGGCTT TGCCCAATCA TAATCAAGGG	2220
AGCGACCGAC TGGAAATGCAA CTCCAAGAAC GACTGGGAGT CCAATCCCAA AGTATTTGTT	2280
GAGTTGGAGT TGGAGGAAGG TTGCCACCCCC ACACATGAAG ATATCTGTAG AAATCAGGTA	2340
GGTCAACTGC TCAGCTGAAT AGCCAAGGGC TGTCGAATC ATGATGGAA CCAGGATAGA	2400
TCCTGAGTAC ATGGCTAGTA AGTGTGCAA GCCAAGAACG GCTGCTTGCG AGTGTGTTTC	2460
TTGAGTTGC ATTAGAGATC TGCCCTCTTA AATACGACTT GACCATTTTC AAAACAATCC	2520
AAACGAGCAA GTGATAGGAC AGGGTAGCCT GCTTTTCAA GCAAATCAGC ACCATCTTGG	2580
AAGGATTTCT CAATCACGAT ACCGATAGCT TGGACTGTGG CACCGGCCTG TTCGATGATT	2640
TGAATCAAGC CTTTAGCAGC TTGGCCATTA GCAAGGAAAT CGTCGATAAT CAAACCTTG	2700
TCCTCTGGTG AGAGGAATTTC TTCAGCGATA GAAACGGTGC TGTCACCTG CTTGGTAAAG	2760
GAGTAGACTT GAGCAGTTAA GATGCCTTCG TTCATGGTGA TGTTCTTAGC TTTTTGGCG	2820
AAAATCATGG GAACTTAA GGCTTCAGCT GTAAAAACGG CTGGGGCAAT ACCCGACGCT	2880
TCAATGGTTA CGACCTTGCT AATGCCAGTA GTAGCAAATT TTTCCGCAAA AACCTTACCA	2940

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ATCTCTCGCA TCAAGCTAAA GTCAACTTGG TGGGTTAAAA AGGAATCTAC CTTGAGGATG	3000
TTATCACCCA AGATATGCCC ATCCTTGAGG ATGCGCTCTT CTAATAATTT CATAAGACCT	3060
CCTAAAGTCT AAAAGTTAAT TTACTTGTG TTTAAATATT TCTATAGTGA TCCCTTTGCT	3120
TAATACTATA TATTGATAA AACTATTACG AGCGAACGGA GTCTTATCAA ATATTTCCCG	3180
TTGTAGTGGT ATCATAGACA ATAATCTGT TATTGTCAT GACGGGATTT TTGAGAGTAA	3240
AATAGTCGG GGAACATATT TAGCCTAACGC CTAGAAATGA AAGAGCTAGG GGCTCAAAAA	3300
TTAGGGATGA AATCCCTGG ATTCCTGAAA TTATTACAG GATAATTCA CCTCCCGTCC	3360
GCACTAATTA AGGAAATAT TAAAAAAAGA CCTACTTAAT CTCTAAGTAA GTCCCCCTAAA	3420
TAGACATGGC AAAAACGCC ATATCTCACT GCTGACTTAC TTATTGTTAG GTGTTCCGGC	3480
ACCTTGTAGA AACGTCGTGC CAATTACAGA CATAAACAG TAAAACGATA TTCAATTAA	3540
AATAGGCTTG AGCCAATGTT TTTATTTAC ACTAAATAAC TTTAGAAATC AACTATTTG	3600
TTAGTGTGTTT GGTTAAAAA ACGAACAAAA AGAAGAGAGG GTGAACAAAA ACTCCATTGT	3660
AAGCTAACAG TTATACTAAA TGAAAATCAA AGAGCAAAC AGGAAGCTAT CCACAACCTC	3720
AAAACACTGT TTTGAGGTTG TGGATAGAAT TGACAGAGCC AGTATCATAT ACCTACGGTA	3780
AGGCGACGTT GACGTGGCTT GAAGAGATTT TCGAAGAGTA TTAGAAGATT TTCCCATCAT	3840
AAAAGGCATA CTATCAAGCT TTTAGACACC TGACAATATG CCTTTTCTA ACTTTAAAGA	3900
CTTTTCCCAA TTTTATTAT TCTACTCGCT AAATCTTAA AAATAGCCAT CTGGATCCAA	3960
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TATGCCAAAG GAGAGATTGA CTCCACGACC AAAGGGATAG GTCAGTTAG CTAGTTGATC	4140
CTTTGTTCCC TCCTCTAACAA TTAGTTGACA CTCTTCAAGA GAAAGAGAAA GTTTCTTCT	4200
GGACGTTGGT ATTCAATCCT AAAACCCAGT AAACCACAGT AGAAGGACCG GGACTGTTCG	4260
ATATTGATA CAAGCAACTC GGGAAATGACC GCATTGAGT CCATATAGAA AATCCTTACA	4320
AGTCAATTTC CAAGACAATC GGTGTATGGT CTTGGCGAGC ACCTGAGTCA ATCATATCAG	4380
ATTTAGTGAC CTTGTCAGCG ATACGGTTAC TTGTGAGCCA GTAGTCGATT CTCCAGCCTG	4440
TATTGTTGAT TTTAGAAGTT TTGCTGCGTT GTGCCACCA AGTGTAGCGT TCAGGAACAT	4500
CGCCATGAAC ATGGCGGAAG GTGTCTGTAA ATCCAGTGC CAAAGGTTG GTAAATCCAG	4560
CACGTTCCCTC GTCAGTAAAT CCAGGTGAAC GGCGGTTGCT AGCAGGATTT GCAAGGTCGA	4620
TTTCATTGTC GGCTACGTTG TAGTCACCGG TCGCAAGGAC TGGTTTTCT TTGTCTAGTT	4680

570	
CAGCCAAATA CTCAGCATAT TTGGCATCCC AGACCTGGCG TTCTTCCAAG CGTTTGAGAC	4740
CGTCACCAGC GTTTGGAGTG TAAACTGGG TTACGAAAAA TGCACTAAAT TCTAGAGTGA	4800
TGATACGACC TTCCAAGTCC ATGGTAGAAG GGGCACCGAT TTCTGGGAAG CTGATAGTAG	4860
GTGTAAGTTC TTTCTTATAA AGGAACATGG TTCCAGCATA GCCTTACGG GCAGGCTCTT	4920
GGGAAGAGCG CCACGTGTT TCAGTGCCTG GGAAGAGTTC TTCTAAAATT TCCACGTGTT	4980
TCTTTGTTAGG TCCTTGGCA GAAAGCTTGG TTTCTTGGAT AGCAATGATA TCAGCATT	5040
CAGCGACCAA GGTTTGTAGG ACTTCTTGGG ACAATTGGC ACGAGCTGAG TCACTAGTTA	5100
GGGCAGCGTT TAGGAATCA ATATTCCATG AGATAAGTTT CATAAAAGTTA CCTTTTTCAT	5160
TCAGATTATA GATTTTATTA TACCAAAAAA AGATCTATTT CCCAACGTA TGGTTTGAAA	5220
AATTACTCTC TTTCGTTAT AATTAAGAAT GATTTTATGA AAGGGAGTGA AAATACATGA	5280
AATTCTACTC TTATGACTAT GTACTCAGCC AAATCGGTCA GCAAAATGGT ATCATGGTTG	5340
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CTTTTGGTCA GCATCAGCAG TTATCAAAAC AATCAAGTTT CTAACAATAA ATTTCAAAC	5520
TCACCTCATT TCATCGAGGT TGTTTCCAAA GATTTGTGAG TAGACAAGTC AGAAAGTCTAT	5580
GTAAATACCTT CCACAAACAC AGATGGCGCA CTTATCAAGG TGGGAGATCG CTATTATCGT	5640
GCCCTAAATG GAAGTGAGCC AGACAAGTAC CTGTTAGAGA AAGTCGAATT GTATAAGACA	5700
GACGCAATTG AACTGGTGGA TGTGACACAA TGACACTTAA TTATATCGAA ATTTCATCA	5760
AACTGGCTT GACTCTCAAA TAGCTCAACA ACAATGTTCA CTTTGTGAAA CGTTTGATTG	5820
ATGGTAAGCC AACTCTCTT ATCAAAATG GGAATATTGA CCCAGAAGCC TGTCGTTCA	5880
TTGGTTTGTC TGCACTGGAT GTATCCCTCA AACTTCGTAG CCAAGGGATT TTCCAGATGA	5940
AGCAAGTCAA ACAGCTGTG CAAGAGCAA ATGGGCAACT CATCGTTGTG CAAATGGGAG	6000
ATGAAAATCC TAAGTATCCA GTTGTGACTG ACGGTGTGAT TCAAGTAGAT GTCTTGAAT	6060
CGATTGGTCG TAGCGAAGAG TGGTGCTTG ATAACCTCA TAAACAAGGG CATGACAATG	6120
TAGCCAATAT CTTTATTGCT GAATATGACA AGGGTGTGT TACAGTCGTAA CTTTATGAAT	6180
AAGAAAAACC TGGGGTCTTG TACTCTCGA AAATCTCTTC AAACCGCGTC AACGTCGCCT	6240
TGCCGTATGT AGGTTACTGA CTTCGTCAGT TCTATCTACA ACCTCAAAGC AGTGCCTTGA	6300
GCAGCCTGCG GCTAGTTCC TAGTTTGCTC TTGATTTTC ATTGAGTATT GGCCTCAGGT	6360
TTCCATTGCA AATCAGAAAG GGATTTATG TCCATTATTC AAAACTTTG GTGGTTTTC	6420
AAGTTAGAAA AACGCCGTTA TCTAGTCGGA ATTGTGGCCC TGATCTTGGT TTCCGTCCTC	6480

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AATCTCATTC CTCCTATGGT TATGGGGCGG GTCAATTGATG CCATCACATC GGGGCAATTA	6540
ACCCAGCAGG ACCTCCTTCT TAGCCTATTT TACTTGCTAC TTGCAGCCTT TGGTATGTAC	6600
TATTTGCGCT ATGTGTGGCG TATGTATATC CTTGGGACCT CTTATTGCTT GGGACAGATC	6660
ATGCCGGTCTC GCTMPGTTAA GCATTCACA AAAATGTCGT CAGCCTTTA TCAAACCTAT	6720
CGGACGGGTG ATCTGATGGC ACACGCAACC AATGATATCA ATGCCTTGAC TCGTTTAGCA	6780
GGTGGCGGTG TCATGCTGC GGTGGATGCC TCTATCACGG CTCTGGTGAC TTTGTTGACC	6840
ATGCTCTTTA GCATCTCATG GCAGATGACT CTTGTTGCCA TTCTCCCCCT ACCTTTCATG	6900
GCCTATAACGA CTAGTCGCCT AGGGAGAAAG ACTCATAAGG CCTTTGGCGA ATCCCAAGCT	6960
GCTTTTCTG AACTCAATAA CAAGGTACAG GAGTCCGTAT CAGGTATCAA AGTGACCAAG	7020
TCTTTCGGTT ATCAGGCAGA CGAGTTGAAG TCTTTTCAGG CAGTCAATGA ATTAACCTTC	7080
CAAAAGAACCG TGCAAACCAT GAAATATGAT AGTCTCTTG ACCCTATGGT TCTCTTGT	7140
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ATTACAGTTG GGAATCTAGT CACCTTTATC AGCTATTGAG ATATGCTGGT CTGGCCTCTT	7260
CTGGCCATCG GTTTCCTCTT TAATACTACT CAGCGAGGGA AGGTTTCTTA CCAGCGGATT	7320
GAAAATCTTT TGTCTCAGGA ATCTCCTGTA CAAGACCTG AGTTTCCTCT GGATGGTATT	7380
GAAAATGGGC GTTTGGAGTA TGCCATTGAC AGCTTGGCTT TTGAAAATGA GGAAACACTG	7440
ACGGATATTC ACTTTAGTTT GGCAAAAGGG CAAACACTGG GCTTGGTTGG GCAGACAGGC	7500
TCTGGAAAA CGTCCTTAAT CAAGCTCCTC TTGCGTGAAT ACGATGTGGA TAAGGGTGCC	7560
ATTTATCTAA ACGGTCACGA TATTGGGAC TATCGCTGA CAGACCTTCG CAGTCTCATG	7620
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GGCAATCTA ACTTGCCCTT TTCAAGCGGTG CAGGAAGCTA CTAAGCTAGC CCGGGTTAC	7740
CAAGATATTG TAGACATGCC TCAAGGATTT GATACTGCTGA TTGGTAAAA AGGAGTCACT	7800
CTTTCTGGTG GTCAAAAGCA ACGGTTGGCT ATGAGTCGGG CTATGATTT AGACCCGTGAT	7860
ATCTTGATTT TGGATGATTC CTTATCCGCC GTAGATGCCA AGACAGAGTA TGGCATTATC	7920
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GCTGTTGTCC ATGCAGATTT TATTTTAGTT CTACAAAATG GTCAAATTAT CGAACGAGGC	8040
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TTGGAAATGA AAGGAGAAGA AGATGCAGAA TAAACAAGAA CAATGGACTG TATTGAAGCG	8160
CTTGATGTCT TATCTCAAGC CTTATGGACT CCTGACCTTT TTGGCACTCA GTTTTCTCCT	8220

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AGCGACGACG GTCATTAAGA GTGTCACTT CCTCGTGGCT TCCCACCTTA TCGACCAAGTA	8280
TCTCAGGAAT CTTAACCAAC TAGCCGTTAC CGTTTGCTG GTCTACTATG GTCTCTACAT	8340
CCTACAAACT GTAGTTCACT ATGTCGGCAA TCTTCTCTT GCGCGCGTGT CTTACAGTAT	8400
TGTTAGGGAT ATTGCGCGG ATGCCCTTGC CAATATGGAG AAACCTGGCA TGTCTTACTT	8460
TGACAAGACG CCAGCAGGTT CTATCGTTTC TCGTTTGACC AACGATAACCG AGACGATTAG	8520
TGATATGTTT TCTGGGATTT TATCCAGCTT TATCTCAGCA GTTTTATCT TTCTGACAAAC	8580
CCTTTATACC ATGTTGGTGC TGGATTTCG TTTGACGGCT TTAGTCTTGC TCTTCTTCC	8640
TTTGATTTTC CTTTGGTCA ATCTCTATCG AAAAAGTC A GTGAAAATCA TCGAGAAAAC	8700
CAGAAGTCTC TTGTCAGATA TCAATAGTAA GCTGGCAGAG AATATCGAGG GAATCAGGAT	8760
TATTCAAGGCC TTTAATCAAG AGAAGGCCCT GCAGGCAGAA TTTGATGAAA TCAACCAAGA	8820
ACACTTGGTC TACGCCAACC GTTCTGTAGC CTTGGATGCC CTCTTTTGA GACCTGCCAT	8880
GAGTTTGTG AAACCTCTAG GCTATGCAGT CTTGATGCCCT TACTTTGGCT ACCGTGGTTT	8940
TTCTATCGGG ATAACGGTCG GGACCATGTA TGCCCTTATC CAGTACATCA ACCGCCTTTT	9000
TGACCCCTTG ATTGAGGTGA CGCAAAACTT TTCAACTCTG CAAACGGCTA TGGTTTCTGC	9060
AGGTCGTGTC TTTGCCCTGA TAGACGAGAG GACCTATGAA CCTCTTCAAG AAAATGGCA	9120
AGCCAAAGTC CAAGAAGGC ATATCCGTT TGAACATGTG TGTTTCTCAT ATGACGGTAA	9180
ACATCCGATT CTGGATGACA TTTCTTCTC TGTTAATAAG GGTGAAACCA TTGCTTTCT	9240
AGGTCACTACA GGTCAGGG AATCGTCTAT TATCAATGTC CTCATGCCCT TTTATGAATT	9300
CCAGTCAGGG AGAGTCTCT TGGATGATGT GGATATCAGG GATTTCTGTC AAGAAGAGCT	9360
GAGAAAAAAC ATCGGTTGG TCTTGCAGGA ACCCTTCCTC TATCATGGAA CTATTAAGTC	9420
CAATATCGCC ATGTACCAAG AAACCAAGTGA TGAGCAGGTT CAGGCTGCCG CAGCCTTTGT	9480
GGATGCAGAT TCCTTTATTC AAGAACTTCC TCAGGGGTAC GACTCCCTG TTTCCGAGCG	9540
TGGTTCGAGC TTCTCTACTG GGCAACGCCA GCTTCTTGC TTTGCTAGAA CAGTCGCCAG	9600
CCAGCCTTCT ACTATTAAG ATGCCAACTG CATCTATGTC TTGGATAAGG GACGCATTAT	9660
CGAGAGTGGAA ACCCATGAGG AACTCTTGGC TCTGGGAGGA ACCTATCACAA AGATGTATAG	9840
TTTGCAGGCA GGGGCCATGG CCGATACTCT TTGAAATCT CTTAAACCA TGTCAAGCTT	9900
ATCTGCAATC TCAAAGCTGT ACTTTGATTT TCATTGAGTA CTAGAAGGAA ATCCTTCAA	9960
TTACAGATTT CTTTCACCGC CTTTCCATT TTGTGGTATA ATGAAAATG TTGACAAATA	10020

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GTATAATAAA AACAAAGGAG AACAGCATGC TGAAATGGGA AGACTTGCCT GTGGAAATGA	10080
AATCAAGCGA GGTTGAGTCT TACTACCAGC TTGTCCTCAA AAGGAAGGGT TCGCTGATT	10140
TCAAGCGTTG CTTGGACTGG GTTTGGCCT TGGTCTTACT GGTCTGACC TCTCCCATCT	10200
TTCTCATCTT GAGCATTGAG ATCAAGTTGG ATAGCAAAGG GCCAGTGATT TACAAGCAAG	10260
AGCGTGTGAC CCAGTACAAC CGTCGGTCA AGATTGGAA GTTCCGTACC ATGGTGACGG	10320
ATGCGGATAA AAAAGGAAGT CTGGTACTT CTGCTAACGA TAGCCGCATT ACCAAGGTTG	10380
GAAATTTCAT CCGACGTGTC CGTTGGACG AACTGCCTCA GTTGGTCAAT GTCCCTAAAG	10440
GTGAGATGTC CTTGTCGGT ACACGACCTG AAGTGCCACG TTATACAGAG CAGTATAGCC	10500
CTGAAATGAT GGCAACCTTG CTCTTGCAAG CAGGGATTAC CTCTCCAGCC AGCATCAACT	10560
ACAAGGATGA GGACACAATT ATCACTCAAA TGACGGAGAA AGGTCTGTCA GTTGATCAGG	10620
CCTATGTGGA GCATGTTCTT CCTGAAAAGA TCGCCTATAA CCTCGCCTAT CTCCGAGAGT	10680
TTAGTTCTT TGGGGACATC AAAATCATGT TTCAAAACCGT GTTGGAGGTA CTAAAATAAA	10740
GTAGTCATAA GAAAATGAGT ACAGATAAAA GGAGCAAATC AATGCCAAAT TACAATATT	10800
CATTTTCACC GCCTGATATC ACAGAAGCAG AAATTACTGA AGTAGTGGAT ACCCTGCGTT	10860
CTGGTTGGAT CACAACAGGT CCTAAAACAA AAGAACTGGA GCGCCGCTTG TCTCTTTACA	10920
CACAGACACC TAAGACTGTT TGTCCTCAACT CTGCGACAGC CGCTCTGGAG TTGATTTTAC	10980
GCGTTTGGGA AGTGGGACCT GGTGATGAAG TCATCGTTCC AGCCATGACC TATAACGGCTT	11040
CATGTAGTGT CATTACGCAC GTGGGAGCAA CCCCTGTATC GGTGGATATC CAAGCAGATA	11100
CGTTTGAGAT GGACTATGAC CTGCTTGAGC AAGCTATCAC TGAGAAAAGT AAGGTGATTA	11160
TTCCAGTAGA GCTCCCAGGG ATTGTTTGCG ATTATGACCG TTTGTTCCAA GTCGTGGAGA	11220
AAAAACGTGA CTTCTTTACC GCTTCAAGCA AGTGGCAAAA GGCCTTTAAC CGTATTGTCA	11280
TTGTCCTGA TAGTCCCCAC GCTTTGGAT CTATTTATAA AGGACAACCT TCTGGTTCTA	11340
TCGCTGACTT TACTTCCTTC TCATTCATG CAGTTAAGAA CTTTACAACG GCAGAAGGTG	11400
GAAGTCCGAC TTGGAAAGCC AATCCAGTGA TTGATGACGA AGAGATGTAC AAGGAATTCC	11460
AAATCCTTTC CCTTCACGGG CAAACTAAGG ATGCTCTTGC CAAGATGCAA CTGGGGTCAT	11520
GGGAATACGA TATCGTTACA CCAGCCTATA AGTGAACAT GACCGATATC ATGGCTTCAC	11580
TTGGTTTGGT ACAATTGGAC CGCTATCCAA GTTTGTTGCA ACGCCGTAAG GACATMTGTGG	11640
ACCGCTATGA TAGTGGTTT GCAGGTTCTC GCATCCATCC TTTGGCACAC AAGACTGAAA	11700
CTGTCGAATC TTCACGCCAC CTCTACATCA CCCGTGTAGA AGGAGCAAGC CTAGAAGAAC	11760

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GCAACCTCAT	CATCCAAGAA	TTGGCTAAAG	CAGGAATTGC	AAGTAATGTT	CACTACAAAC	11820
CGCTTCCTCT	CTTGACAGCC	TATAAGAAC	TTGGATTG	TATGACGAAC	TATCCTAAGG	11880
CCTATGCCCT	CTTGTGAGAAT	GAAATTACCC	TCCCTCTCA	TACTAAATT	AGCGATGAAG	11940
AAAGTAGACTA	TATCATTGAG	ACTTTCAAAA	CAGTTCTGA	AAAAGTGCTA	ACTTTATCAA	12000
AAAAATGACA	AACTACAGTC	AAGCGAAAGT	GATCCTGCC	CTAAAAAGTC	TAATTGAGTG	12060
TAAAAACTGT	TGTTTCAAT	TGATAATAGT	TTACACCTGT	AGTTGAGGCC	CCTTTCTCCT	12120
CAGAGAGAGA	ATTTTTATAG	GATTTCCCTT	TCCTGTGGGA	GTCCCGTGGT	TTGAAATAAG	12180
ATGTGAGCAA	TTTGTGTTAG	CATTAGAAT	CCTTACTAGA	CATCATTTAG	AAAATCTAGT	12240
GTCTTGTCT	AGTTTCAAT	TCACCCATT	TTTGAAAGA	CGTGAGTTTC	CATGAGTGAG	12300
ATTGTGGAAA	CTCGCGTCTT	TTTTGTGTTT	CAGAATATTG	TTCAAAATTT	TGTGCCTGTC	12360
TTTCATGTTTC	TAGTCATTCT	TTTGCGATGAT	AGAATTTATA	GCATGTTGAT	ATTATAATAA	12420
TACAAATATT	CTATATGTTT	AGTGATGCTT	GCTATACATT	ATTAGATCTC	CTGCGAGACA	12480
ATCTATAAAA	CACTGTCTA	CGATTACCTA	TATGCCCTAT	TCCAGTATTT	TAGAAGCACT	12540
GCATCTATT	TTATCGAGGT	AAATCTAGC	TTTTATAGAA	GGTCTATTTA	AGAAATATAT	12600
TGTTAGTGT	TAGTTCAAT	CCGCCATATG	AGCGATATTG	AGGTAAATAT	CCCTGGCGAA	12660
TGCTTGTATG	ACAAGGTATT	TGTTCTTCA	TTTATAATTT	ACAACATATC	AAACAAATTTA	12720
AATATAGTAA	ATGGGATATT	TTATATTCAA	GCTAAGAAAG	ATAGCATCAC	TTTTGAATGG	12780
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GTAGATATAG	AAAAATGAAA	TGAGAATAGG	ACAAATTGAT	CGGGACAGTC	AAATCGATT	12900
CTAACAAATGT	TTTAAAGTA	GAGGTGTACT	ATTTTAGTTT	CAGTCTACTA	TAGAACTGAC	12960
CAAGTCAGTA	ACCTAGACTT	AGGGCAAGGC	GGCACTGACC	TAGTTGAAG	AGATTCCGA	13020
AGAGTATAAA	TTTTAATATT	TTCTTGTGTT	ATTCCTTGAC	AATTCAATTT	GGAAAATATA	13080
TGATAAAAGAT	AATGACAGCG	GTGTCATTCT	ATCTATTTA	AGAAAAGTAA	TAATCAATTG	13140
TTAAAAATAG	AAAAAAATT	GGAGGTTCTG	ATGAAATATT	TTGTTCCG		13188

(2) INFORMATION FOR SEQ ID NO: 71:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 32768 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 71:

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AACGAGTGCA	TCAGTCTCAG	CAAGCACCAAG	TGGCTCGGCC	TCAGCAAGCA	CCAGCGCGTC	60
TGAATCCGCA	TCAACCAGTG	CCTCAGCTTC	AGCAAGTACC	TCAGCATCTG	AATCAGCATC	120
AACAAGTGCA	TCGGCTTCAG	CAAGCACAAAG	TGCTTCAGCC	TCAGCAAGTA	TCTCAGCGTC	180
TGAATCGGCA	TCAACGAGTG	CGTCCGCTTC	AGCAAGTACT	AGGCCTCAG	CATCAGCGTC	240
AACAAGTGCT	TCGGCTTCAG	CGTCAACGAG	TGGTCTGAG	TCAGCATCAA	CGAGTACGTC	300
AGCCTCAGCA	AGCACATCAG	CTTCTGAATC	TGCATCAACC	AGTGCCTCAG	CCTCAGCATC	360
GACAAGCGCC	TCAGCTTCAG	CAAGTACCAAG	TGGCTCAGCC	TCAGCAAGTA	CCAGTGCCTTC	420
AGCCTCAGCG	TCGACAAGTG	CGTCGGCCTC	AACAGTGCA	TCTGAATCGG	CATCAACCAG	480
TGGCTCAGCC	TCAGCAAGTA	CTAGCGCCTC	AGCCTCAGCA	TCAACGAGTG	CGTCCGCTTC	540
AGCAAGTACT	AGTGCATCAG	CATCAGCATC	AACGAGTGCA	TCGGCTTCAG	CAAGTACCAAG	600
CGCCTCAGCT	TCAGCAAGCA	CCAGTGCCTC	AGCCTCAGCA	AGTACCAAGCG	CCTCAGCCTC	660
AGCAAGCACC	AGTGCCTCAG	CTTCAGCAAG	TACCAAGTGC	TCAGCCTCAG	CGTCGACAAG	720
TGGCTCGGCT	TCAGCAAGTA	CCTCAGCGTC	TGAATCAGCA	TCAACGAGTG	CATCAGCTTC	780
AGCATCAACA	AGTGCCTCAG	CTTCAGCAAG	TATCTCAGCG	TCTGAATCGG	CATCAACGAG	840
TGGCTCCGCT	TCAGCAAGTA	CTAGCGCCTC	AGCCTCAGCG	TCAACAAGTG	CTTCAGCCTTC	900
AGCCTCAACG	AGTGCCTCAG	AGTCAGCATC	AACGAGTAG	TCAGCCTCAG	CAAGCACATC	960
AGCTTCTGAA	TCTGCATCAA	CCAGTGCCTC	AGCCTCAGCA	TCGACAAGCG	CCTCAGCCTTC	1020
AGCAAGTACC	AGTGCCTCAG	CCTCAGCAAG	TACCAAGTGT	TCAGCCTCAG	CGTCGACAAG	1080
TGGCTCGGCC	TCAACCAGTG	CATCTGAATC	GGCATCAACC	AGTGCCTCAG	CCTCAGCAAG	1140
TACTAGCGCC	TCAGCCTCAG	CATCAACGAG	TGGCTCCGCT	TCAGCAAGTA	CTAGTGCATC	1200
AGCATCAGCA	TCAACGAGTG	CATCGCTTC	AGCAAGTACC	AGGCCTCAG	CTTCAGCAAG	1260
CACCAAGTGC	TCAGNCTCAG	CAAGTACCAAG	CGCCTCAGCC	TCAGCAAGCA	CCAGTGCCTC	1320
AGCTTCAGCA	AGTACCAAGTG	CGTCAAGCCTC	AGCGTCGACA	AGTGCCTCAG	CTTCAGCAAG	1380
TACCTCAGCG	TCTGAATCAG	CATCAACGAG	TGCATCAGCT	TCAGCATCAA	CAAGTGCCTTC	1440
AGCTTCAGCA	AGTACCAAGTG	CGTCCGCTTC	AGCATCAACG	AGTGCCTCAG	TCTCAGCGTC	1500
AACCAGTGCC	TCTGAATCAG	CATCAACAAG	TGCTCGGCT	TCAGCAAGCA	CCAGTGCCTC	1560
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AACGAGTGCT	TCGGCTTCAG	CATCAACGAG	TGGCTCAGCC	TCAGCAAGCA	CATCAGCTTC	1680
TGAATCTGCA	TCAACCAGTG	CGTCCGCTTC	AGCGTCAACC	AGTGCCTCAG	CTTCAGCGTC	1740

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GACAAGTGCT	TCGGCTTCAG	CATCAACGAG	TGCGTCCGCC	TCAGCAAGCG	CAAGTACCTC	1800
AGCGTCAGct	TCGGCCTCAA	CCAGTGCCTC	GGCTTCAGCA	AGCACAAGTG	CGTCAGCCTC	1860
AGCAAGTATC	TCAGCGCTCG	AATCGGCATC	AACGAGTGCG	TCTGAGTCAG	CATCAACGAG	1920
TACGTCACTC	TCAGCAAGCA	CATCAGCTTC	TGAATCTGCA	TCAACCAGTG	CGTCAGCCTC	1980
AGCATCGACA	AGCGCCTCAG	CTTCAGCAAG	TACCAAGTGCT	TCAGCCTCAG	CGTCGACAAG	2040
TGCGTCCGCC	TCAACCAGTG	CATCTGAATC	GGCATCAACC	AGTGCCTCAG	CCTCAGCAAG	2100
TACTAGTGCA	TCAGCCTCAG	CATCAACGAG	TGCATCGGCT	TCAGCATCAA	CCAGTGCCTC	2160
GGCTTCAGCG	TCAACCAGTG	CGTCAGCTTC	AGCAAGTACCG	AGTGCCTCAG	TCTCAGCCTC	2220
AACAAGTGCT	TCAGCCTCAG	CATCGACAAG	TGCCCTCGGCT	TCAGCAAGCA	CATCAGCCTC	2280
TGAATCAGCG	TCAACCAGTG	CTTCAGCCTC	AGCAAGTACCG	AGTGCCTCAG	CTTCAGCCTC	2340
AACCAGGCC	TCGGCCTCAG	CAAGCACCTC	AGCTTCTGAA	TGCCCTCAA	CCAGCCTC	2400
GGCCTCAGCA	AGCACCTCAG	CTTCTGAATC	GGCCTCAACC	AGCAGCTCAG	CCTCAGCCTC	2460
AACGAGTGCT	TCGGCCTCAG	CAAGCACAAG	CGCCCTCGGT	TCAGCATCAA	CGAGTACGTC	2520
AGCTTCAGCG	TCAACCAGTG	CTTCAGCCTC	AGCATCAACA	AGTGCCTCAG	CCTCAGCAAG	2580
TATCTCAGCG	TCTGAATCGG	CATCAACGAG	TGCCTCTGAG	TCAGCATCAA	CGAGTACGTC	2640
AGCCTCAGCA	AGCACCTCAG	CTTCTGAATC	GGCCTCAACC	AGTGCCTCAG	CCTCAGCCTC	2700
GACAAGGCC	TCAGCCTCAG	CAAGTACCGAG	TGCCCTCAGCC	TCAGCGTCAA	CAAGTGCCTC	2760
GGCCTCAGCA	AGTGCATCTG	AATCGGCATC	AACCAAGTGCG	TCAGCCTCAG	CAAGTACTAG	2820
TGCATCGGCT	TCAGCATCAA	CCAGTGCCTC	GGCTTCAGCG	TCAACCAGTG	CGTCAGCTTC	2880
AGCAAGTACCG	AGTGCCTCAG	TCTCAGCCTC	AACAAGTGCT	TCAGCCTCAG	CATCGACAAG	2940
TGCCCTCGGCT	TCAGCAAGCA	CATCAGCCTC	TGAATCAGCG	TGCACAAGCG	CCTCAGCTTC	3000
AGCAAGTACCG	AGTGCCTCAG	CCTCAGCCTC	GACAAGTGCG	TCAGCCTCAG	CAAGTACTAG	3060
TGCATCAGCT	TCAGCATCAA	CGAGTGCCTC	GGCTTCAGCG	TCAACCAGTG	CATCAGAGTC	3120
AGCAAGTACCG	AGTGCCTCAG	CTTCAGCCTC	AACAAGTGCG	TGCCCTCAG	CAAGCACCAG	3180
TGCCCTCGGCT	TCAGCAAGTA	CTAGCGCCTC	AGCCTCAGCC	TCAACCAGTG	CGTCAGCCTC	3240
AGCAAGTACCG	TCAGCGCTCG	AATCGGCATC	AACGAGTGCG	TCCGCTTCAG	CAAGTACTAG	3300
GGCCTCAGCC	TCAGCGTCAA	CAAGTGCCTC	GGCTTCAGCG	TCAACGAGTG	CGTCAGCTTC	3360
GGCATCAACG	AGTGCCTCAG	CTTCAGCAAG	TACTAGCGCC	TCAGCCTCAG	CGTCAGCAAG	3420
TGCATCGGCT	TCAGCATCAA	CGAGTGCCTC	CGCTTCAGCA	AGTACTAGCG	CCTCAGCCTC	3480
AGCGTCAACA	AGTGCATCGG	CTTCAGCGTC	AACGAGTGCG	TCTGAGTCAG	CATCAACGAG	3540

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TGCGTCGGCT TCAGCAAGTA CCAGTGCCTC AGCCTCAGCA AGTACCAAGTG CGTCAGCCTC	3720
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AGCCTCAGCA AGTACTAGTG CATCAGCTTC AGCATCAACG AGTGCATCGG CTTCAGCATT	3840
AACCAGTGCA TCAGAGTCAG CAAGTACCAAG TGCGTCAGCT TCCGCATCAA CAAGTGCCTC	3900
GGCTTCAGCA AGTACTAGCG CCTCAGCCTC AGCGTCAACA AGTGCCTCAG CTTCCGCTC	3960
AACCAGGCC TCGGCCTCAG CAAGTATCTC AGCGTCTGAA TCGGCATCAA CAAGTGCCTC	4020
GGCTTCAGCA TCAACCGAGTG CATCAGTCTC AGCAAGCACC AGTGCCTCGG CCTCAGCAAG	4080
CACCAGCGCG TCTGAATCCG CATCAACCAAG TGCGTCAGCT TCAGCAAGTA CCTCAGCATT	4140
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ATTTTACTG GCTTCTCTCA TAGCGATTCC AAAATGACT GCCCAAGATA AGATTCTAAT	30180
ATAGTTAGCA GTAAGCAGGG CGTTGACTGG GTGTCAACC AGTTGAGCA AGAGGTTGCT	30240
GAGAACCTGC CCAATCCCCT CTGGTGGTGC AATTCAGTA TTGGCACTAT TTGGGTAAT	30300
TTCAATAGGG ACGATGAAAT TTGCTAGTAC AGCTACAAGA GCAGCGGCCA AAGTCCCTAT	30360
CATAGGATAT ACAAGAAAAC AACAGTTTC ATATTGCTAT CTTGTCCTT TTGATGTTGG	30420
GAAAGGGCAT TGGCAACGAG AGCAAAGACT AGGATAGGAG CAACAGCTTT TAGACCTCCA	30480
ACGAATAAA CCTCGAGTAG CCCAATCCCT GAGAGATTAG GAAGGGTCAG TCCTAGGATT	30540
CCCCACAAAG CATACCAATC AAGATACGCT TGACAAGGCT TGCCATTATTC CAAGCATGAA	30600
TGATTCTTT CATAATAATC TCCTTTTGT GTAGTGATTA TGATTATAGT ATAAATGATA	30660
GACAAAATCA AGAATTTCAGT GTCTATTTTG TGAATATTAA TGGAGAATGA GACTGATGAA	30720
AATATGGTAT AATGAAATAA AGGAGTTTA TATGAAAAAA TTTTATTCAAG CTTATATTGA	30780
AAAGCTAGAT GTGACAACCA TTATCGAGAA TATTCTAACC AAGGTCATTT CTCTTTACT	30840
GCTTTTAATT GTATTTTATA TTGCTAAAAA AATGCTTCAT ACCATGGTGC AGAGAATTGT	30900
CAAACCTTCT CTAAAAATGT CTCGTCATGA TGTTGGACGC CAAAAAACCA TCTCACGTTT	30960
ACTAGAAAAAT GTGTTTAATT ATACGCTATA TTTCTTTTA CTCTACTGCA TTTTGTGAT	31020
TTTAGGTTTG CCAGTTCTA GTTGCTGGC TGGAGCTGGT ATTGCTGGGG TAGCGATTGG	31080
TATGGGAGCC CAAGGCTTTC TGTCTGATGT CATCAATGGC TTTTCATCC TCTTTGAACG	31140
TCAACTGGAT GTGGGAGATG AGGTCGTTCT GACAAATGGA CCGATTACTG TATCGGGTAA	31200
GGTTGTCAGT GTGGGAATTC GTACGACACA GCTTCGTAGC GAGGAGCAAG CCCTTCACCTT	31260
TGTCCTAAC CGAAATATCA CAGTTGTTAG CAATTCTCA CGCACAGACT AGACCTGTTA	31320
TTTTAAGTAA TTTGTGGTAC AATAGAGGGA GTTAAATAAG GAGAAAAGAT GGTTTAGAA	31380
AAGCAGTTGG GCAATGGTTG TACCTGGATA GACCTAGACC TAGGAAAGTT GAATAAACTA	31440
GAAGACCTTT CTGAAATTAA CGGTTGGAC AAGGAAACCA TTGAATACGC ACTGGATAGA	31500
AAAGGAGCGCG CCCACATGGA CTACCCACGT GAAAGTGAGA CGGTTACCTT TATCTATAAT	31560
GTCTTAGACG TAAAAAAAGGA CAAGGCCTAC TATGAGACTT TTCCCATGAC CTTTATTGTC	31620
GAGCAGTCGC GCCTGATTAC CATTAGTAAT ACCAAGAACG CCTATGTCA TGAACAGATG	31680
ACTCGTTATC TGGAGAACCA TGACACGCTT TCGATTATA AGTTTCTCTT TGCCAGTCTG	31740
GAAATCATCA GCAATGCCA CTATCCTGTC ATTGAGCAGA TGGACAAGAG TAGGGATGAG	31800
GTCAATGACC TCTTGCGCCA GCGAACTACC AAGAAAAACC TCTTGTCCCT GTCTGATTG	31860

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GAGACTGGTA TGGTTTATCT GACGGCAGCT GCCAAACAAA ATCGGATTGTT GTTAGAGCAT	31920
ATTCAAGGTC ATGCCTTGTA TCGTAGTTTT GATGAGATTG AGAGAGAACAA GTTTGATGAT	31980
GCCATGATTG AGGCTCATCA GCTGGTATCC ATGACAGACC TAATCTCTCA GATTTTACAG	32040
CAGCTTCAG CCTCTTACAA CAATATTCTA AACATAATC TGAATGACAA TTTGACAACC	32100
TTGACTATCA TTTCAGTCTT GCTAGCTGTT TTGGCAGTCG TGACAGGCTT TTTCGGAATG	32160
AATGTTCCCT TACCTTTAAC AGATGAGCCC CATGCTTGGC TCTATATCAG TTTGGCTAGT	32220
GCAGGTTTGT GGATTGTTT ATCCTTGTAA CTAAGGAAAA TTGCGAAAAA AAGTTAAGAA	32280
AAGGAGCCAG AATGGCGATT GAAAATTATA TACCAAGATTT TGCTGTGGAA GCAGTCTATG	32340
ATCTGACAGT CCCAAGCCTG CAGGCGCAGG GAATAAAAGC TGTTTGGTC GATTTGGATA	32400
ATACCCCTCAT TGCTTGGAAC ACCCTGATG GAACGCCAGA GATGAAGCAA TGGCTACATG	32460
ACCTTCGGGA CGCGGGTATT GGCATTATCG TAGTGTCAAA TAACACCAAA AAACGCGTTC	32520
AACGAGCAGT TGAGAAAATTG GGGATTGATT ACCTTACTG GGCCTTGAAG CCCTTCACAT	32580
TTGGTATTGA CCGTGCTATG AAGGAATTCC ACTATGACAA AAAGGAAGTG GTCATGGTTG	32640
GTGACCAACT CATGACAGAT ATACGAGCAG CCCACCGTGC AGGGATTCGG TCAATTTAG	32700
TCAAACCCCTT GGTCCAACAT GACTCAATCA AAACGCAGAT TAACCGAACT CGTGAGCGTC	32760
GTTGTTATG	32768

(2) INFORMATION FOR SEQ ID NO: 72:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 14872 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 72:

CCAGTCACAA AGAAATTGAG CGCGTTCAGC TGAGGATGCA CTATGATGCA AGCTACATT	60
CATTTGATGG GATATTAAGA AAGGAGATTT TCATGACACT TTTAGATGTA AAACACGTT	120
AAAAAAATTAA TAAAACACGT TTTCAGGGCA ACCAAGTAGA AGCCCTCAAG GATATTCACT	180
TTACCGTAGA AAAGGGTGAC TACGTTGCCA TCATGGGTGA GTCTGGTTCT GGTAAATCAA	240
CTCTTCTCAA TATTCTAGCT ATGTTGGATA AACCAAGTCG TGGTCAGGTT TACTTGAATG	300
GAACTGACAC CGCAACTATT AAAAATTCAAC AGGCTTCTAG TTTCCGGCGT GAAAAGCTAG	360
GATTTGTCTT CCAAGACTTT AACCTGCTAG ATACTCTGTC TGTAAAGGAC AATATCTTGC	420

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TTCCGCTTGT CTTGTCAAGA AGACCTATAA CGGAGATGAT GAAGAAATTG GTGGTGACAG	480
CTGAGAACATCT GGGTATTAAC CAATTGCAAG AGAAGATCACC TTACGAGATT TCTGGTGGTC	540
AGAAAACAGCG TGTAGCAGTA GCCCGGCCA TCATCACAGA ACCTGAATT CTCCCTGGG	600
ACGAGCCAAC AGGAGCCCTT GATTCCAAGT CATCTGCAGC CTTACTTGAT GTCTTTAATG	660
AAATCAATGA GCGTGGCAA ACCATCCTCA TGGTAACCCA CTCAACAGCA GCTGCTAGCA	720
GGGCCAAGCG TGTTCTCTT ATCAAAGACG GCATTCTTTA CAACCAAATC TACCGTGGAG	780
AGAAGACAGA GCGTCAGATG TTCCAAGAAA TCTCTGATAC CTTGACTGTC ATGGCAAGCG	840
AGGTGAATTA GTATGTTTCG ATTAACCAAT AAGTTAGCGG TATCGAACTT GATTAACAAAC	900
CGCAAACACTCT ACTATCCCTT TGCACGGCT GTTCTCTTGG CAGTCACCAT CACCTATCTC	960
TTTTACTCCC TAACCTTCAA TCCAAAGATT GCGGAATCC GTGGAGGAAC CACCATTCAA	1020
GCAACACTTG GATTGGTAT GTTGTGCGTT ACCCTTGGT CACCATATTAC GTCTCTATG	1080
CCAATAGTTT TGTCATGAAA AACCGTTCCA AGGAACGGG TATATATGGC ATGTTAGGCT	1140
TGGAGAAGCG CCATCTAACAT AGTATGACCT TTAAGGAGTT AGTGGTATTT GGGATTCTAA	1200
CTGTTGGAGC GGGTATCGGT ATTGGAGCCT GTTGTGACAA GTTAATTTC GCTTCCCTGC	1260
TCAAACATAAT GAAACTGAAG GTTGAGCTGG TTGCTACCTT CAAATGAAT GTTGTCAATTG	1320
CAGTACTTGT TGTCCTTGGA TTGATTTCC TAGGCCTCAT GTTCCCTGAAT GCTCTTCGAA	1380
TCGCCCGTAT GAATGCCCTC CAGCTCTCGC GTGAGAAAGC AAGCGGAGAG AAAAGAGGTC	1440
GCTTCCTTACCC TCTCCAAACG ATTCTTGGTT CCATAAGTTT AGGGATTGGC TATTATCTTG	1500
CCCTTACGGT AACCGATCCT CTTACAGCCC TAACAACTTT CTTCCCTAGCT GTTTGCTGG	1560
TTATCTTGG TACTTATCTA TTGTTTAATG CAGGGATTAC AGTCTCCCTA CAAATCTAA	1620
AGAAAAACAA GAAATACTAT TACCAACCTA ATAACCTCAT ATCTGTTCC AACTTGATT	1680
TCCGTATGAA GAAAATGCG GTTGGACTAG CAACCATCGC TATTTGTCA ACAATGGTTT	1740
TGGTAACCAT GTCAGCAGCG ACAAGCATT TCAATTCCGC AGAAAGCTTT AAAAAAGTTC	1800
TAAATCCTCA TGATTTGGG GTTTCAGGGC AAAATGTTGA AAAAGAAGAT TTGGACAAAC	1860
TCTTGAGCCA GTTTGCAAGT GACAAAGGTT ATAGTGTCAA AGAGAAAGAA GTACTTCGTT	1920
ACAGTAACCTT TGGTATTGCA AATCAAGAAG GAACCAAGTT AACTATTTTT GAAAAAGGAC	1980
AAAACCGTGT CCAACCCACCA ACAGTTTCA TGGTATTGAA CAAAAAGAT TATGAAANTA	2040
TGACTGGTCA AAAACTGTCT CTATCAGGAA ATGAGGTCGG TCTCTTGCC AAAATGACG	2100
GACTGAAAGG ACAGAAAGCT CTAACCTCAA ATGATCATCA ATTTCTGTC AAAGAAGAAT	2160
TTAATAAAGA TTTCATTGTG AACCAGTTC CAAATAAGTT TAATATCTTG ACTACTGATT	2220

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ACAATTACCT	TGTTGTCCT	GATTTACAAG	CCTTTTGGA	TCATTCCCA	GATTCGGCTA	2280
TCTATAATCA	GTTTACGGT	GGTATGAATG	AAATGTCAG	TGAAGAAGAA	CAACTCAAGG	2340
TCGCTGAGGA	GTATGAAAAC	TACCTCAATC	AATTAAATGC	TCAATTAGAC	ACAGAAGGTA	2400
GCTATGTTA	TGGTAGCAAT	CTAGCAGATG	CTAGTCTCA	GATGAGTGCC	CTCTTGTTG	2460
GTGTCTCTT	TATCGGTATT	TTCCATATCCA	TTATCTTTAT	GGTCGGAAC	GTTCTGGTCA	2520
TCTACTACAA	ACAAATTCT	GAAGGCTACG	AAAGACCGTA	ACGCTTTATT	ATCTTGAGA	2580
AAGTCGGTTT	GGACCAAAAG	CAAATCAAGC	AAACCATCAA	CAAACAGGTT	TTAACTGTTT	2640
TCTTCCTTCC	TTTGCTCTT	GCCTTCATAC	ATCTCGCCTT	TGCCTACCAT	ATGCTTAGCC	2700
TGATTTAAA	AGTGATTGGT	GTACTGGATA	CGACTATGAT	GTTGATTGTG	ACCTTGTCTA	2760
TCTGCCCTAT	CTTCCTCATC	GCCTATGTGC	TGATTTTCAT	GATTACTTCA	AGAAGTTATC	2820
GCAAGATTGT	GCAAATGTA	AAAAGATACC	TCGACTTCAA	AATCGAGGTA	TTTCTTGTAT	2880
TCTAAATGCT	GAAAAGTTGT	CCGAGCAGGA	AGGTAACCTC	CATGGTCAAG	AGACCAATAG	2940
CAAGGTTCCG	AATCATAGCT	TTTMTGGTTG	GGGCTTTCC	AAAGTCTAGCA	CTTGTGTAAC	3000
CAGTGAGAAG	AAGGGCCACA	CCGACAATAA	GGACGGTAGC	AGGGATGCGG	TAATCACTTG	3060
GAAAAATGGT	CACTGACAGC	ATTGGAGGCA	AACTTCTAAG	GAAAAGGCA	ACGAAGCTAG	3120
AAATGGCAGC	GTGCCAAGGA	TTGGTAAATT	CTTCATAC	AATCCCATAT	TTTCTCTCTA	3180
CCAGAGCCTT	GAGTGGATTT	TTAAGAAAGA	TCTTATTGGT	CAAGAGTTGG	GCAGAAGTTT	3240
TGAATTCTCC	ATTTTGGATA	TAAGCAGCAT	AGAGGGATTT	TTTGGCTAGT	TCCCTATCTT	3300
GGTCTAGCAA	GAGTTTTCT	CGCGAACCG	CAGCTTCCTC	GGTATCTTTT	GGAGTTGAAA	3360
CGGATACATA	TTCTCCACCA	GCCATTGAAA	AGGCACCAGC	TAAGATAGCC	GTAAACCTG	3420
ATAAAAAGAT	AATCCAGATA	TTGGTCGTGG	CACTGGCAAC	TCCGATAACC	ACACCAGCAA	3480
TGGAATAAT	TCCATCGTTA	GCATCAAGAA	CACCCGCACG	CAGGATATTT	AAACGACCTG	3540
CAAAATTTGA	ATCAATTTCG	TGATTTGT	CTGACGCTAA	ATTTCAAGTT	CAAGTTAGCC	3600
ATCAAGAAGT	CTTCTCTGGG	TGACTTGTAG	TCCAAGCATT	TTTTAGGATA	GTTGTTAAC	3660
CACTTTTCGA	TGAATGCGAC	TTCTTGGGA	GTCATTTCT	TGGTCCCTT	AGGTAACCAT	3720
CTACGAATGA	GCCTGTTGTG	ATTCTCATTA	GTCCCCCTT	CCCAAGAGGC	ATAGGGATGT	3780
GCATAATAAA	TGTGCTCCTC	AGAAAATACA	TTAGACAAGC	GATTGAATT	CGTTCCATT	3840
TCTGCCGTGA	TGGAAAAGAAT	CTTGTGTTGT	TTTAAGATGA	GTTTAGAGC	CTGATTGACC	3900
ACATCAGCAC	TTTATTGGA	AATCAATCGG	ATGATCTGAT	GTCTACTTTT	TCGATCCGTC	3960

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AAGACAAGCA	ACGAGTAGTT	TTTCGCTCTC	GTAAGTAGAA	CTGTATCAAT	CTCATATAATGC	4020
CCATTCTCCA	AGCGAAAATT	GATAGCTTCA	AGCCGCTGTT	CGATGGATTG	ACCAGCAGGT	4080
TTAAAGTTGG	TGCTGCCCTG	TTTCTTAAGC	GCTTTTCCTT	TTCTAGGGTA	AAGCAGATCC	4140
TGTTTGCTTA	ACCCCAATT	TCCATGATGA	ATCCAATAGT	AAATGGTTGA	AATTCCCACG	4200
TTAACCCCTT	TAGCCATCAC	CATCATTTCA	GGCGAAAATT	TTTGGTTATG	ATAGTGGAGA	4260
ATCTTTCTT	TTAGTTCCCTT	GGTCAAGCTT	GATTCTTGA	CCGAGCGCTT	GCGATTGTTT	4320
TCATAAGACT	GTTGAGCATA	GTCGGCAGAA	TAAACCTCTT	TGAAGCGCCC	TTTTCCAAGA	4380
CATTGTCGGA	CTGTCCCCACG	CTTGATTTC	GTGTGGATAG	TTTGAGGAAC	TTTTCCAAGC	4440
AGAGAGGCAA	TTTCTCTATT	TGATTTCCCT	TCTTTTTCC	ATCTTCGAT	TAAGCGACGG	4500
CTATCGATTG	TCAAATGTT	GCCTTTGTA	GTATAATGGT	TTTGATCTC	TGTGCCCTTC	4560
TTGTGTTTGT	GGTTGAACAA	CAAGTATAAC	ACAGAGGTGT	TTTCTTATGC	CTACAAGAGC	4620
TATCGGCTAG	TTGAACCATC	TAATTTTAG	GAGGGCTGGG	TGGCTAACTT	CATTATAGAA	4680
CTTCATTTA	CGAACATATA	GTAAAATGAA	ACAAGAACAG	AACAAATCGA	TCAGGACAGT	4740
AAAATCTATT	TCTAACATG	TTTTAGAAGC	AGAGGTGTAC	TATTCTAGTT	TCATCTATT	4800
ATATTTTGT	TTTTTATCAA	AAAATACTTT	ACAAGTTCTT	AAAACATGA	TATAGTAATA	4860
AAGCTTAGAA	AATGAGATGA	TGTTTTCTAG	CAAATATAAA	CCCGAGTAAA	AAATGCCTAC	4920
GGACAGGCAG	GGTTGAATGC	CGAACGCTGG	TTGAAAAGCC	ACATTATTGA	TAGGGTTAAA	4980
AGCCTACTTT	TATAAGTTGA	TGTTAGGACA	CTTGTCTAA	TTCATAAATT	TTTAGTGTGG	5040
TGAAAGCACA	CGTCATCTTG	TGAAACGATC	AATAAAGTAC	GTAATATTG	CTACTAGAGA	5100
GTTAGGAAAC	ATCGGGAAACA	GACATACTCA	ACAGAAACCA	AAATAAACAC	GTCAGAACAT	5160
TGCAGAGCAG	GTGAAAACCT	GCTCTTTT	CATGAGTCAA	CCTTTAGTTTC	CTTAGTTTC	5220
ATAAGGTCT	AAAATATTG	AAAGGAGTAT	GTGTTGAAAG	AGTTAGATCA	AAACCAAGCC	5280
CCAATTATG	AGGCCTTGGT	GAAGTTACGC	AAGAAAAGGA	TTGTTCCCTT	TGATGTTCCA	5340
GGTCACAAGC	GTGGACGGGG	AAATCCAGAA	CTTGTGAAAC	TCTTAGGAGA	AAAATGTGTA	5400
GGCATTGATG	TCAATTGAT	GAAACCTTTG	GATAATTAG	GCCATCCTAT	TTGATTATT	5460
CGTGATGCG	AGGAGCTGGC	TGCAGATGCT	TTGGAGCTA	GCCATGCTT	TCTAATGATT	5520
GGTGGAAACAA	CTTCATCGGT	GCAGACTATG	ATTCTGGCAA	CCTGCAAGGC	AGGAGATAAG	5580
ATTATTCTGC	CACGAAATGT	CCATAAAATCT	GCTATCAATG	CGTTGGTTCT	ATGTGGTGCC	5640
ATTCCCATCT	ATATCGAGAT	GAGTGTAGAT	CCTAAGATTG	GTATCGCTTT	AGGTCTTGAA	5700
AATGACCGAG	TAGCACAGGC	CATAAAGGAC	CATCCAGATG	CTAAGGCTAT	CCTAACAAAC	5760

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AATCCTACTT ACTACGGCAT CTGTTCAGAC CTAAGGGGT TGACAGAAAT GGCTCATGAA	5820
GCTGGCATGA TGGTTTAGT AGATGAAGCC CACGGAGCGC ATTTGCATTT CACTGATAAA	5880
CTTCCAATTG CTGCTATGGA TGCAAGGGCT GATATGGCAG CAGTTCCAT GCATAAGTCT	5940
GGTGGGAGTT TGACCCAAAG CTCCATTGGA CTTATCGGGG ACCAGATGAA TTCTGAATAC	6000
GTTCGTCAGA TAATTAACCT GACCCAGTCT ACATCTGCCCT CTTACTTGTT GATGGCTAGT	6060
TTGGATATTT CACGTCGCAA CTTGGCCCTT CGTGGTAAAG AGTCGTTGAA GAAAGTCATT	6120
GAGCTATCTG AGTATGCCCG CGGTAAATC AATGCTATCG GTGGCTACTA TGCCTACTCA	6180
AAAGAGTTAA TAGACGGTGT TTGGGTTGACG GATTTTGACG TAACTAAGCT GTCAGTTAC	6240
ACTCAGGGTA TTGGCTTAAC AGGTATCGAG GTTTATGACC TCTTGGGAGA CGAATACGAC	6300
ATTCAGATCG AGTTGGTGA TATCGGCAAT ATCTTGGCCT ATATTTCCAT CGGCGACCGC	6360
ATCCAAGACA TCGAGCGCTT GGTTGGTGCT CTGGCTGATA TTAAGAGACT CTATTCAAGA	6420
GATGGAAAAG ATTTGATAGC AGGAGAATAT ATTCAAGCCCG AGTTAGTGCT GTCTCCGCAA	6480
GAAGCCTTCT ATTCAAGAAAG AAAAGTTA ACCTTGGATG ATTCTGTTGG ACAGGTCTGT	6540
GGAGAAATTG TTATGTGTTA CCCTCCAGGT ATTCCCTATCT TGGCTCCTGG TGAACCCATT	6600
ACACGAGAAA TTGTCGACTA TATCCAATTC GCCAAGGAAC GTGGTTGCTC CCTCCAAGGG	6660
ACCGAAGATC CAGAGGTCAA TCATATCAAC GTTATTAAGA GAAAGACAAA CTATAAGAAA	6720
ACTCAATAGT TTATCTAAA CTATTTCTTTA TTCAATTGATG GATGATTTA GATGATTTA	6780
GAGCACGGCA AAAAGCCCTT GAATTAGAAG CGGTCAATCG CTTAATTCT ATCAGCTTAT	6840
CAAATCCCTGC CTCAAGCCTT TTCTGAGGAT TAGGGTAGCG TGTCAAGAGT TGGTAGGTAT	6900
ATTCTGAATG CTTTCCAACG ATTTTATCCA ACTCAGGAAA GATGATATCA AGACAACGAG	6960
TGTATTGTAC TTTCCAATCA GACTGTTTT TCTTGAGACG ATGAATATGT CTAGCCAGTA	7020
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GTTTAAAGAGC GATGCCATGA GCGCTTTCT TATCCGTTT AGTTTGCAGA AGTGATAATG	7140
ATTTGGCAAA TTCTTGATG AGCAAAGGAT TGTAGGTGTA AACTTATAT CCTTGTTCAT	7200
GCAGGAAGTT CAGTAGATTA AAGGCATAAT GTCCGGTATT TTCAAGAGCG ATGAGACAGT	7260
CTTGGTTGAG CTGTCGAAGA GACAGATCTA AGAGTTCAAA ACCAGCTTTA TTATTTGAAA	7320
AAGTGAGTGG TTTAAGAACAA GTTTTCCCTG GAACATTCAA GGCTGTAACA TCGTGTTAT	7380
TTTTAGCGAC ATCAATGCC ACATAAAGCA TGGGAGTATC TCCAGATATA GTATTTCAAG	7440
TCTACTGGGT TATCCACGAA CTTTTGCGCT TGGTACCTTA GACGAGATAA AACGTCTATG	7500

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CGTTATCAA CTCATTACCA ATTGAAACAA AAAACTGTGG TTAGAGCCTT TCGGAAATCG	7560
TCAAGCGATT GGAGGAAATG AACTAATCCA CAGTGGCTTA TTCCAAGTAT ACCACTTGGG	7620
CTTTGGCAGT AGCTAACTGC GCTAAATATA ATATAAGGAG AAATAGATGG ATTTATGGTT	7680
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TGACGGCGGT GTTGCCCAAG TATTAACCCCT CTATCCTGAA CTGGAGCAAA TTGATATTGT	7980
GGAACCGGAT GAGATGTTGG TCGAGGTCTG TCGTGAGTAT TTCCCAGACT TTGCTGCAGG	8040
GCTAGATGAT CCTCGTGT TA CCATTTACTA CCAAAATGGG CTACGCTTT TGCGAAACTG	8100
CGAAGATGAT TACGATATTA TCATCAACGA TGCGACAGAT CCATTTGGCC ATACGGAAGG	8160
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GATTTACCAAG CATGGGAGTC CCTTCTTGA CGAGGATGAG TCGGCCTGCC GAAGCATGCA	8280
CCGCAAGGTC AATCAAGCCT TTCCAATCAG TCGGGTTTAT CAGGCCATA TTCCAACTAG	8340
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CGTGGGAGCC TTTATGTTGC CCAAGTATGT TGAGGACATT TTAGAAGAAG AGGAAGGAAA	8520
AAAATGAGTC GTTTACTAGT TATTGGTTGT GGGGGCGTTG CCCAAGTGT TATTTCAAAG	8580
ATTTGTCAAG ATAGCGAAC ATTACAGAG ATTATGATTG CTAGCCGTAC CAAGTCAAA	8640
TGCGATGACT TGAAAGCGAA GCTAGAAGGC AAAACAGTA CTAAAATTGA AACTGCAGCA	8700
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CGTGCTATCT ACGAAAAACG TTGTAAGGAA CTTGGTTTA CAGCCTACTT TGACTACTCA	8940
TGGCACTGGG CTTATCAAGA GAAATTCAA GAAGCAGGCT TGACTGCTCT TCTTGGTTCT	9000
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ACCAACTTTA ATCCAGAAAAT TAATCTCCGT GAGGTTCTG CGCCAGGTT TCAGTGGAA	9180
GATGGGAAAT GGGTCGAAGT CGAAGCTATG TCTATCAAGC GTGAGTATGA TTTCCCTCAA	9240
GTTGGACAAA AAGATATGTA TCTCCTTCAC CATGAAGAAA TCGAATCATT GGCAAGAAC	9300

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ATTCCAGGTG TCAAACGCAT TCGTTTCTTT ATGACTTTG GTCAATCTTA CTTGACGCAC	9360
ATGAAATGTC TTGAAAATGT TGGACTCCTT CGTACGGATA CCATTAACCT TAACGGCAA	9420
GAAATTGTC CAATTCAATT TTTGAAAGCC TTGCTTCCAG ATCCTGCCAG TCTTGGCCA	9480
CGTACAGTCG GAAAAACCAA TATTGGATGT ATCTTTACAG GTGTCAAAGA CGGTGTCAA	9540
AAGACTATCT ATATCTACAA TGTCTGCGAC CATCAGGAAT GTTACGCAGA GGTTGGTCG	9600
CAAGCTATTT CTTATACGAC AGGAGTTCCA GCCATGATTG GGACAAAATT AGTCATGAAC	9660
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TAATGAAGTT AGAACAAAGT CCAACACCAG CCTATGTTAT TGACTTGGCC AAGTTAGAAG	9840
CTAATTGCCG CATTCTACAA TATGTACAAG AAAGAGGCCGG TTGCAAGGTC TTGCTTGC	9900
AGAAGGCATA TTCCCTCTAC AAAACTTATC CCTTGATTAG CCAGTATCTA TCAGGTACGA	9960
CAGCTAGTGG ACTCTATGAG GCCAAATTGG CAAGGGAAGA ATTTCCCTGGT GAAGTCCATG	10020
TATTTGCGCC TGCTTCAAG GATGCAGACT TGGAGGAATT GCTAGAGATA ATGGACCATA	10080
TAGTCTTAA CTCAGAGAGA CAGTTGCGTA AACACGGTCC GCGTTGTCGA GAGGCTGGT	10140
TCAGTGTGG TTTGCGCCTC AACCCCTCAGT GTTCAACTCA AGGcAGATCA CGCGCTCTAT	10200
GACCCTTGTG CACCAGGTT TCAGCTTGGTGA GTTACTATAG ACAAGATTCC GAGTGATTTG	10260
CTAGATTTGG TTGACGGACT TCATTTTCAT ACCCTTGGG AGCAGGGAGC AGATGATT	10320
CAAACAACTT TGAAAGCAGT AGAAGAACAG TTTGGTCCCT ACTTACATGA GGTAAAATGG	10380
CTCAATATGG GTGGTGGTCA TCATATTTACA AGAGAAGTTT ACGATGTGGA TTTGCTGATT	10440
TCAGAAATCA AGCGTATCCG AAAAACTTAC AATCTTAAA TCTATATCGA GCCTGGTGAA	10500
GCCATTGCGC TTAATGCGGG TTATTTAGCA ACTGAGGTAT TAGATATTGT AGAAAACGGT	10560
ATGGAAATCT TGGTTTACA CGCCTCTGCG ACCTGCCATA TGCCTGATGT ACTTGAGATG	10620
CCCTATCGTC CACCTTTGAG AAATGGCTTT GAGTCACAGG AAAAAGGCCA TACCTACAGA	10680
CTTTCTCTA ATACCTGTCT GACGGGGCAT GTGATTGGTG ATTATAGTT TGAAAATCCA	10740
GTCCAAATCG GAGACAGACT TTATTTCAAA GACATGGCCA TTTATTCTTT TGTCAAAAT	10800
AATACCTTA ATGGTATTGG ATTGCCAAGT CTCTATCTCA TGGACGAACA GGGAGACTGT	10860
AGCTTACTCA AAGCTTTGG CTATCAAGAC TTTAAAGGGA GATTATCATG ATGGACAGTC	10920
CAAAAAAATT AGGCTATCAC ATGCCAGCAG AGTACGAACC CCATCATGGT ACCCTCATGA	10980
TATGGCCGAC TCGACCAGGA TCATGGCCTT TTCAAGGAAA GGCTGCTAAA AGAGCATTAA	11040

600

CTCAGATTAT CGAGACCATA GCAGAAGGG AAAGAGTCTA TCTTTGGTG GAGCAGGCCT	11100
ATCTATCTGA AGCCAATCC TATCTGGAG ACAAGGTTGT TTATTTAGAC ATTCCCACCA	11160
ATGATGCCTG GGCGCGTGAT ACTGGCCCAA CCATTCTCGT CAATGATAAA GGTAAGAAAT	11220
TAGCCGTGGA TTGGGCCTTC AATGCTTGGG GAGGCACCTA TGATGGTCTT TATCAAGATT	11280
ATGAAGAGGA TGACCAAGTA GCCAGTCGTT TTGCTGAGGC CTTGGAAAGG CCTGTCTATG	11340
ATGCTAAACC TTTGTACTG GAAGGAGGCG CAATCCATAG CGATGGTCAA GGAACATTTC	11400
TCGTAACTGA AAGTGCTTG CTTAGTCCTG GTCGCAATCC TAACTTGACT AAAGAGGAGA	11460
TTGAAAACAC ATTATTAGAA AGTCTTGGTG CTGAAAAAGT TATTTGGCTT CCTTATGGTA	11520
TTTATCAGGA TGAAACCAAT GAACACGTCG ATAATGTTGC TGCCCTTGTGTT GGTCCCTGCTG	11580
AGCTTGTGTTT GGCTTGGACA GATGACGAAA ATGATCCCCA GTATGCCATG TCAAAAGCAG	11640
ATCTCGAACT CTTAGAACAG GAACACAGATG CAAAGGTTG TCACITCACC ATTCATAAAT	11700
TGCCTATCCC TGCAGTTGTC CAAGTTGTGA CAGAAGAAGA TTTGCCAGGC TACATCTATG	11760
AAGAAGGAGA AGAAAAGCGA TACGCAGGTG AACGACTAGC AGCTTCCTAC GTAAACTTTT	11820
ATATGCCAA CAAGGCTGTC TTGGTTCCAC AGTTGAGGA TGTAACGAC CAAGTGGCCT	11880
TAGATATCCT CAGCAAGTGT TTCCCAGACC GTAAAGTTGT CGGAATACCA GCCAGAGATA	11940
TTCTCTTAGG TGGTGGCAAT ATCCACTGTA TCACCCAACA AATTCCAGAA TAGGAGAAA	12000
AGATGAGAAA TGTAAGAGTT GCAACCATTC AGATGCAATG CGCTAAGGAT GTGGCAACAA	12060
ATATCCAAAC CGCAGAGCGT TTAGTACGTC AGGCTGCTGA GCAAGGAGCC CAAATTATTC	12120
TCTTGGCCGA GTTGTGAA CATCCCTATT TCTGTCAGGA ACGTCAGTAT GACTACTACC	12180
AGTATGCCCA ATCTGTAGCG GAAAATACTG CCATTCAAGCA TTTTAAGGTG ATTGCTAAGG	12240
AACTACAAGT TGTTTACCA ATCAGTTCT ATGAAAAGA TGGTAATGTC TTGTATAACT	12300
CTATTGCCGT CATTGATGCA GATGGGAAAG TGCTGGCGT TTATCGAAAG ACCCATATAC	12360
CAGATGACCA TTATTATCAA GAAAATTCT ATTCACGCC TGGTAACACT GGTTCAAGG	12420
TCTGGAATAC TCGCTATGCT AAGATTGGTA TCGGTATCTG TTGGGATCAA TGGTTCCCTG	12480
AAACAGCGCG CTGCTTGCA TTGAATGGTG CTGAATTGCT CTTTATCCT ACAGCTATCG	12540
GTTCAGAGCC AATTTGGAT ACAGATAGTT GTGGTCACTG GCAACGTACT ATGCAAGGGC	12600
ACGCAGCAGC GAATATTGTT CCAGTCATCG CAGCCANTCG TTATGGTTA GAGGAGGTTA	12660
CTCCTAGTGA GGAAAATGGC GGACAGAGCT CCAGTCTTGA CTTCTACGGT TCCTCCTTTA	12720
TGACGGATGA AACAGGAGCT ATTCTAGAAC GAGCTGAAAG ACAAGAAGAA GCTGTTCTGT	12780
TAGCTACTTA TGACCTAGAC AAGGGAGCAA GTGAACGCCCT AAACCTGGGGC TTGTTTCGAG	12840

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ATAGAAGACC AGAAATGTAT AGACAAATTA CAGATTAGTG TGGGAGAAAT GAGAGATTCA	12900
TTCTGCTAGA CTAACCTCTT ATTAGTAACT ATAAGATACT ATGGCATCTA GTAAATCGAT	12960
TTTATGATT CGCTATTCTT GTCTATTGAT TAGTCCGTAT TTTAAATAT TAGCAAAAAA	13020
GCAAATAGCA GTAACCTCTG TCTATTTGCT TTTCTTTTT ATAGAATATA TTTCTCAATA	13080
GCACCGCAA CGCCCTCTTC TTCTGCTT GAGGTAACGG CATCCGCAAG AGATTTGATA	13140
TAATCGCTGG CATTCCCAT TGCAATCCA AGCCCTGCAA ACTGGAGCAT TTGATATCG	13200
TTATTAGCAT CGCCCATGGC CATAATCTCT GAGGAATCAA TCTTCAAAAT CTCAGCTAGT	13260
CGTGAAGAG CAGTAGCCTT TGTCGTTCCA AGCGGCATTG CTTCTAAAT GACAGGCTGC	13320
GAACGAACTC CACTGAATCG TTGGCAAAGC TCTTCAGCAA AACGCTGCTC AAAATCGTCT	13380
GTGGTTCTT TTGTCCTAA ACACATACCT TGGAACATCC GGAACTTTCC ACTAGTCGCT	13440
TCTTCAGAG AAATTTCAGT CAGGTCTGAA AATACTAGTT TAGCATCATT TTCAATAACT	13500
TGATGGGCT TGTCACCGAG AACAAAATAA TGTGACTCGT CAAAAAGTGT CAACTGAACA	13560
TCACTTTT CAGCAAGGTC ATAGAGGTAT TCGATGTCAG CTGGACTCAG TTCTTCCAG	13620
TCAACTAGAC TCCAATCACT GGTCTGGTGA GPTGAACAAAC CGTTGTTAAC AATAATATAT	13680
TCGTTCTGGA GGTCAAGCTC CAGTTTTTG TAGTAGGGG GGACACCGAA AAGGGGGCGA	13740
CCCGTACAGA GAACCACTT GACACCTTT TCAATGGCTT TGTGAATAGC AGTAATGTGT	13800
GCTTGTTGGGTTTCCCTGGC TTCATTGAGG AGGGTGCCGT CCATATCCAA GGCTAGTAGT	13860
TTAACATAG GTCTCCCTCT TTATCTTGC TATTATTATA GCATATTTG GAGAAGAAAT	13920
TGATAGAAAG CTTGAGACTA ATTGATTTTA TAGTTTAAGA TGTTTGATG ACAATTTCATG	13980
ATTTGAAGAG GATATTCGC AAAGATATGC TATACTATGT TTGTCATGT TGCAACTAGA	14040
CAAATAAAA AACCAACTTA ATATAATGT TTTTTGTAA GTAGGTATGA GTAGCAGATT	14100
ACTCAACTAA TCTGAAGAAT AATGGAGGAA ATATATCATG ATTTTAATGA CAAAAAATAT	14160
AAATCTAACAA AATGAAGAAT TAGAGCTGAT ACAAGGTGGA CGACATCCAT ATGGAAAAAA	14220
TCCTAATGGT AGGTACGATT GGGAAATAGA ACCAGTATTA ACTCTGCTGG TTGATGGATT	14280
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AACTGCTGCG AGATTTAAG TAAATTTAT TAGGAATATG AAGAAACAAG GGGAGAAAAC	14400
AGAGGATTTA ATATGAAAAA ACCAGCTATT CAAATTTAC TAGCATTGTC CTTAATTTT	14460
TACAAATCAA CTTGGTTTG GAGGCTTTTC AATTATCTCG CAAAGCCCTA TCTACCAGCA	14520
AGTCGTGAAT TTTTCAGAT TCTGCTTTTG ATGGAGAGCG GAGTTCTTTT CTTAGCGGTC	14580

602	
ATCTATCTAC TGGTTTTGCG AGGAAAGAAA ATTTTCATT TCAAGTGGCA GCTGAGGTAC	14640
TTCATCTACC TTTTACTGGG CTACATCATT TCATATATGT CTGACTTCCT CTTTCGTAT	14700
TTCATATCCC TGTCTTCAAA TCAGATTCT TTGAATGAAA CGGTAGAAAT GATGGGGAGA	14760
CAGGAGTTCC CTTATGTCTT GCTCATCGTT TGCTTCATCG CCCCTATTGC TGAGGAATTG	14820
ATTTATCGAG GtGTGCTTAT GACAACCTGT TGCAAAAACt CACCTTGGTA CG	14872

(2) INFORMATION FOR SEQ ID NO: 73:

- (i) SEQUENCE CHARACTERISTICS:
- (A) LENGTH: 10223 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 73:

CGTGCTATCG GTCTCAAAAC CAATCTGGTC GCTATGGTCA AATCCAGTTG GAAAATCCAT	60
TCTTCTTGGG GCCATCTGCT GGATTGCCAT CATCCTCAC C ACTCTTGGTA TGCAGACCCCT	120
TATCGCATT TTCTAATACT CTTCGAAAAT CTCTTCAAC ACAGTCAACG TCGCCTTGCC	180
GTTAGGTATAT GTTACTGACT TCGTCAGTTC TATCTGCAAC CTCAAAACGG TGTTTGAGCT	240
GACTTCGTCA GTTCTATCTG CAACCTCAAAC ACGGTGTGTTT GAGCTGACTT CGTCAGTCGT	300
ATCTACAAACC TCAAAACAGT GTTTTGAGCT GACTTCGTCA GTTCTATCTG CAACCTCAAAC	360
ACAGTGTGTTT GAGCAGCCCCG TGGTAGTTT CCTAGTTGCT TCTTTGATTT TCATTGAGTA	420
TAACACAAAA GGTAGCCCCAT CAGGTACCTT TTCTTATGC TTCCCTCAATC AAGCGAGTAT	480
GTTCTCTCTT GATACAGCGA TTCATCACGA TATCATCACA TCCACCATCA CGCAAAATCT	540
CTTTTCGCTTC TAAACTTTCA AGTCCTAGCT GTGCCAAAAA AATCTTGGCA TCAGCTTTGA	600
GAAAATCACG CGCCACATCG GGCAGAAATT CACTGCGACG ATAAACATTG ACAATATCTA	660
CAGGAAAAGG AATTTCAGCG AGGCTAGCAT AACCCCTTTAC ACCCAAGATT TCGCCACCTG	720
CCGCCTTGGG ATTGACTGGG ATGATTTAT AGCCCCGAGC CTGCATTTC TTTGTTACTC	780
GATTCGCTGGT TGTCTTCA CGGTAGACAA AACCCACAC AGCAAGGGTT TTACTCGTTG	840
CGAGATACTG ACGAACATCG CCATCACTTG GATTGATAAA TTCTTGACTC ATAGAAATCC	900
TCCTTTTCA TCAGTATAGC ACATTTGAA AAGGTTGCA GAATTATACT ACAAAAAGG	960
AGGACTAGCC CCCTTTTAT TTAGCCTCGT ACCAGGTTGC CCCTTCATTC TCATCTGCGA	1020
TAAGAGGAAC ACTGAGTTGA ATGGCTTCTT CCATGGTTG TTTCACCAAT TTTTCATCT	1080
CTACCAATTG AGATTTAGGC ACTTCAAGGA CGATTCATC GTGCACTTGT AACAGCATCT	1140

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TAGTCTGATA ACCACCTGCA ACCAAGGCTT TATCCAGCTG AATCATGGCA ATCTTGAGAA	1200
TATCTGCTGC CGAACCTGG ATAGGTGAGT TGATAGCAGT TCGCTCCGCA AAACCACGAA	1260
TATTGAAGTT GCGCGAATTG ATATCTGGCA ACTCACGGCG ACGCTTAAAG AGGGTCTCTA	1320
CATAGCCCTT ATCACCGGCC TCCCCCACCA CTTCATCCAT GTAGTTTTA ATACCTGGAA	1380
AACGTTCAAA GTAGGTATCA ATGTAGGCTT TGGCTTCCTT ACGACTAATT CCCAAATTAT	1440
TAGACAAGCC AAAGTCTGAA ATCCCATAAA CCACTCCAAA GTTAACTGCC TTGGCATTGC	1500
GACGGTCGTT TGCAGTCACA TCATCAGGAC GCTCAATGCC AAAGACCCGC ATGGCTGTG	1560
AAGTATGGAT ATCTGCCCTC TCTTGGAGG CCTTAATCAA GTGCTCATCC TTAGAAATAT	1620
GCGCCAAAAC GCGCAATTCA ATCTGTGAAT AGTCAGAGCT GAGTAGCACCA CTATCCTCCC	1680
ACTCTGGCAC AAAACCTTC CGAATCAAGC GCCCCTGTT CAATCGGGCA GGAATATTTT	1740
GCAAGTTGG ATCCACACTA GACAAACGCC CGGTCTGGGT CAAATCCTGC ACATAGCGAG	1800
TATGAATCTT TCCATCAGCC AAAATCCAGT CCTGCAAGCC AATTACATAA GTAGATTGAA	1860
TCTTAGCAAT TTGACGGTAA TCCAGGATTT TCTTAACAAAT CGGAGCAATA GGAGCGAGAC	1920
GCTCTAAAAC ATCCACTGCT GTCGAATAAC CTGTCTTGGT TTTCTTAGTG TATTCTAGAG	1980
GAAGTCCCAA TTTCTCAAAG AGAAGCACGC CCAACTGCTT AGGCGAGTTG ACATTAAACT	2040
CCTCACCCAGC CAGCTCGTAA ATCTCTTGAG TCAGTTTTTC AATGACAAGC TCATTTTCAG	2100
CCTGCATCTC AAGCAAGTC TCTTTCTTGA CCATAATCCC AGCAATTTC ACCTTTGGCAA	2160
GGACAAAAGC CAGAGGTTGC TCCATATCAT AAAGAAGCTC TAATTGCCA TTTTCGCTGA	2220
GTTTTCAAG TAAAATAGGC TCTGTTCTA CCAAAACAGC AAGTTTACAA GCTAAGTGT	2280
CCAAGAATTTC CTCACGTTCA GGAATGGCCT TTTTAACACC CTTACCGTAG AAAGTTTCAT	2340
CATCAACCAA GTAAGTCTGA CCATAAAAGAC TAGCGATGGT CGCAATTTC CAATGCTCCA	2400
CAGTCGAAAG GAGGTATTAA GCCAAACGGA TGTCAAAAGC AGGCGCCTGC AAATCCACAC	2460
CAAAACGTTG CAAAAGAACT TTAACCTTCT TAAAGTCATA AACTCTCAGA GATGTTTTT	2520
CTAAGAAATC CTTGAAAATC GGGTCTTGCA ACAGCTCAAG CTTGCTGTG GCATAGAGCT	2580
TATCCCCACA AGACCAGACA AATCCAACCA AATTATCCGT ATGGTAATTC TCACCAAAAA	2640
GCTCAAAGTG GAAGATAGAC TCTTCACTCA GCATATCTTG ACTGATTGG TCAACAATAG	2700
TAAAATCCAA ACTCTCAGAC ACATCAGCTG ACGACACATT TAAAGCCTGC TTTAGCTGTT	2760
TGAAGCCCAT CTCATCGTAG AATTTCCCAA GATTTCAAC ATCTGGACCA CTATAGACCA	2820
AGTCCTCTAA ACCAATCGCA ATCGGTGCCT TGGTATCAAT GGTGCTAGT GTTTTAGACA	2880

604	
AAAAGGCCTG TTCCCTGTCA TTGATGAGAT TTTCCTTCAT CTTAGAAGTC TTCATTCCAT	2940
CAATATTTTC ATAATCCCC TCAAGCGAAC CATGCTCCAG CAAGAGCTTA ATACCCGTCT	3000
TTTCACCGAC TTTGGTCACC CCAGGGATAT TATCCGACTT ATCACCCATG AGGCCCTTGA	3060
GATCGATAAA CTGAGCTGGT GTGAGGCCA TTTCTTCAT GAGGTAATCT GGCCTAAAGG	3120
CCTCAAACTC AGCCACACCT TTCTTGAAA TTTCAACCAC CGTATGCTCA TCCGTAGCT	3180
GAATCAAATC CTTGTCCCCA CTGACAATAG TAATATCAA ACCATCCTGC TCTGCTAGCT	3240
TATCCAGCGT CCCAATGATG TCATCCGCCT CATACTGAGC CAGATCATAG TGACGAATCC	3300
CCATATGATC CAGCAACTCA CGAATGAAAG GAAATTGCTC ACGAAACTCA TCAGGAGTCT	3360
TGGCCCGACC ACCCTTATAG TCCGCATACA TCTCTGTCCG GAAGGTCGTC TTTCCCGCAT	3420
CAAAAGCCAC CAAAATATGA CTCGGCTCAA CCCGCTCCAA TAAATGACTC AACATCAACT	3480
GAAAACCATA AATCGCATTG GTATGAAAC CAGCCACATT CTTAAACGG TCCAACGTCT	3540
GATACAGCGC AAAAACGCC CGAAAAGCTA CAGAAGACCC ATCAATCAAT AATAATTTT	3600
TCTTATCCAT ACACCCATTA TAAAGGAAAG AATCAAAAAA TACCATTGGG AAGAGCTAGA	3660
GCAAGTATTT TTCAAACCTT TTCCGAATAA ATAGATAGAG CCAGAGAAATT TAGTAAACCT	3720
AGATTTAAAA ATGTGCTATA ATATAGTATA TTGAATCTAT AATAGTACAC CTTGACTGCT	3780
AAAAATTTTC TATAAATTAA TTTGACTTTC CTGATAGAGT TATTACATC TTATTTCAAC	3840
TCACTATAGA AGGAGGAATA GGAGGATTCT CAGACATCCG GGCATCAGCC CAACTAATGA	3900
TTTGATTGCT AAGAAAATAT TCAGCAATCC AGAAATCACT TGTCAATTAA TTCGCGATAT	3960
GCTGGACTTG CCAGCAAAAAA ATGTGACCAT TTTGGAGGG AAGCAGATATTC ACGTATTACT	4020
CTCCATGCCT TACTCGGTGC AGGATTTTA TACCACTATA GACGTCTTGG CGGAGTTGGA	4080
TAACGGTACT CAAGTAATTA TTGAGATTCA AGTCCATCAT CAGAATTTT TCATCAATCA	4140
CTTGTGGCT TACCTGTGCA GTCAAGTTAA TCAAAATCTT GAAAAAATTC GTCAGCGAGA	4200
AGGTGATACT CACTAGAGCT ACAAACACAT CGCTCCTGTT TACGCCATTG CTATCGTGG	4260
TAGTAATTAT TTCTCAGATG ACCTGGCTTT TCATAGCTTT AGTATGCGCG AAGACACAAC	4320
AGGTGAGGTA TTGGCAGTTA CCAACAATGG ACAGGAAAC CATCTGGTTA AGATGGCATT	4380
CTTGGAAATTA AAAATACAG AGAAACCAGC AAAGACAAGG TTCGCAAGCC ATGGTTGGAG	4440
TTTTTCCGCA ACAAGCCCTT TACCCAGCAA CGCAGCAGG CCATTACCCA AGCAAATCAA	4500
CTGCTGGACT ACAAGAGCTG GTCCGAGGAG GACAGGAAAA TGTTTAGTCA ACTACATATG	4560
CGAGAAGAAC AAGTCTTGTGTT AGCACAGGAC TATGCCATTGG AACTGCTAG GGCTGAAGGC	4620
CTTGAACAAAG GACTAGAGCG TGGGAAAGTT GAAGGAAGGG CAGAAAGGAA ACTTTTGCC	4680

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TTCCTAGACA TAGTACGCCA AGGTCTCTG ACTTCTGAGG TTGCCAGCCA GCAATTAGGT	4740
ATGTCAGTAT CTGAATTGAG GCACGTGGT TAAAATGGCT CCATAATATC CATACTGGGT	4800
AAATCCCTA TGGATATTAT GGAGCCTATT TTGTGTAGAA AAAAGTCCC ATATGACCTA	4860
TAATGAAAAG CGACAAAACA ACTCATTAGA AAGAACATCA TGGAACAATT ACATTTATC	4920
ACAAAATTAC TAGACATTAA AGACCTAAT GTCCAGATTT TAAACATCAT CAATAAGGAT	4980
ACACACAAAG AAATCATCGC CAAACTGGAC TACGACGCC CATCTGCC TGAGTGCGGA	5040
AACCAATTGA AGAAATATGA CTTCAAAAA CCTTCTAAAA TTCCCTATCT TGAAACGACT	5100
GGTATGCCA CAAGAATTCT CCTTAGAAG CGTCGATTCA ACTGCTATCA CTGTTCAAAA	5160
ATGATGGTCG CTGAAACTTC TGATGACCTA CAGTCATATT TCTTCTCTTT TTATTATATC	5220
ACAGTTTAA ATCTAGCTTT ACTAGATTCA CCGCTACTAT CTATTTATTC GGAAAAAAGA	5280
CGAAAAAAC TGAGAACAT CTCAGGCTTG GTCATTAAT TTTTTCTCA ATATCGAAA	5340
GTGGAGAAAG TGGTCCTTT TCATGAATAC GTACGATAGC ATCCCCTAGG AGATGAGCGA	5400
TTGAAATCTG CTCAATCTTA TCAATCAAAC GCTCTCTGG CAGATAGATG GTATCCAAA	5460
CAACCAATTCTTAATAGCT GATTTTTGGA TATTGTCGGT AGCAGGACCA GAAAGAACTG	5520
GGTGCCTACA GCTTGCATAG ACTTCAACAG CACCAGCTTC CGCAAGAGCA TCTGCCGCAT	5580
GACAAATCGT TCCAGCGGTAA TCAATCATAT CATCAATCAA GATACAAGTC TTGCCCTCA	5640
CCTTACCGAT GATATTCATA ACTTCACTAG TATTTCATCTT ATCAACGCTA CGACGTTTAT	5700
CAATAATAGC GATAGATGTT TTCAAAAATT CTGCCAACTT ACGAGCACGA GTCACCCCTC	5760
CATGGTCCGG GCTGACAACC ACATAGTCAG AACCAACCAT ACCACGACGC TCAAAATAAT	5820
CTGCAATCAG AGGAGCACCC ATCAAATGAT CCACAGGAAT ATCAAAGAAT CCTTGAATT	5880
GCGCAGCATG CAAGTCGATG GTCAATAAAC GATCCACTCC AGCTACTTCA ACCATATTG	5940
CGACAAGTTT TGAAGTGATT GGCTCACGCG CTCTCGCCTT TCTATCCTGA CGTGCATACC	6000
CATAGTAAGG CATGACAACA TTGACAGATT CTGCACTCGC ACGCTTCAAA GCATCTACCA	6060
TAATCAAAT TTCAAGCAGA TTGTCATTTA CAGGCGAACT AGTTGATTGT AAGATAAAGA	6120
CGTGTTCCTTACGGGATTGAT TCTTCAATGT TGACCTGAAT CTCTCCATCT GAAAATTGGC	6180
GAACACTTGA TTCCCCAAC TCTATCCAA TCTCCTGCC CACACGTTCT GCCAATTCTT	6240
TATTAGAAGA AAGGGCAAC AGCTTTAAAT CAGAAAAAGA CATGATTTC TCCGGTATAT	6300
ATGTATAACT TGTGCTTTCA ACAAGATTTT CCATCTACCA TTGTAGCGCT TTTTGCACCA	6360
TTTTCAATC AAAAATAAAAA GAAGGGCACC ATATTGTAC CCTTGCATCA TTCTTTGAA	6420

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AAATATTCTA GGTCACTAAC TCATTGTGTT TCTCAACAAA GCAATAAGCA	TGATAAAAAC	6480
CATAGAGAGC AATAGCCGTA ACCACTGGAA TCGCTAAAGG CAACTCTGTT	TCCAACCTCCA	6540
CAAAGGAGA GTTAAACAAG AAGTGAGTTC CCAAGGCTAA ACCTAGAAAA	ATAAGGCCCT	6600
GTTTCTTGCCT AACCTTCTGT CCTTTATAGG CTCTGTAAAG CAAGTAAACA	CCTACTACAG	6660
CTAGACCTGA AAAAGTCCAG TGAGAGGCAA TTCCGTGAGAT GATACGCTCT	AAAATTCCGCG	6720
AAATAGTAAA GTCAAGCCC TCTGGCAAAT CCGTACGAAT ATAACCAATA	TCCTTAATCA	6780
TTTGGAAATCC CAAACCGGAA GCAATTCCAA GTAAAAACAA AGATTTTAAT	TTTCGCACAG	6840
GAATCAAAGC CAAACAAAAA ACAAGTGACA ATAATTCAA GGGTTCTTCT	ACCAAAGGAG	6900
CCGCAATTAGC ACTTTCAAAG GCATTTAAAA ATGGACTATC TGGGAAAAGA	ACCCCCAGTA	6960
AATCATGGAT ATAAGTATTA GCAAAACTAG ACAACCAGCC TGAAAGGAAC	ATCCCCTCCA	7020
ATAAAAGACAG AATCAAAACC TTCTTTGGCA ATTCCCATTT TTCCCAATAC	GGAAGAGAAA	7080
ATAAAAGAGCC GGAATCATGT AAAAGAGAGC TAGAAAGATA GAAACTCCCA	TTAGTCCATA	7140
TTCCGCACCT GACCTCGAAC CGTCCGTATA GTAGATGGTT TCATACTGTA	AACCAATACA	7200
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AACTATTAT AATTCTACGA CTGTCTACT TCCTGTATCA ACATTGTAAA	TGGCACCGAGA	7320
GATAATGACA TCGTCTGGTA TTAGGGGAGA CTCGATAAGC AGTTGCATAT	CCTCGCGTAC	7380
ACTCTCTTCT ATATCTTGGGAA AGGGCAAGAA GTCTGGTCT GACACATCGA	CACCCAAATTC	7440
TTCCCTCAAA TACTCCTGAA AAGGTTCAATT TTCAAAGGTC TGAGCACCCAC	AGTCTGTATG	7500
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CATATCCTCA GTCACTCGAC CACCTGCATT CCGCAAAATA TGAGCATCCC	CAAGTGCCAA	7620
ACCTAGAGCT TGCGCACCGT GCAAACTGTGA GTCCCATACAG GTCACAATGG	CTACTCTGGT	7680
TTTAGGTTTA AGTGGCAGAT TTAAGTGCCTC ATGTTAGGGCA ACATAAGCCT	GATTGGCTTG	7740
CATAAAACTGT TCAAAATACG ACACGATTCC CTCCCTGAAA ATTTGATAGT	CAAATATTTC	7800
TCCTATCTTA TCATTTTAA GAGAATTGTG CACGGATTAT GCAAAGACCT	TTTTCAAGAC	7860
TTCCGTGAAATC GTTGTACCGC CAATGACCTG AATTTCTTA GGCAGAGTGA	TTCCGTCAA	7920
GGAATTCTTA GGTACATAAA TCTTAGTAAA GCCCAGTTA GCAGCTTCGT	TGATGCCTTG	7980
CTCAATACGA TTCACGCGCC GAATCTCTCC TGTCAAGCCC AGTTCTCCGA	CAAAACATTC	8040
CTGAGGATTA GTTGGCTTGT CTTTGTAGCT CGAAGCAATA GCAACTGCAA	CAGCCAAGTC	8100
AATCGCAGGT TCATCCAATT TAACACCACC AGCAGATTG AGATAGGCAT	CCTGATTTG	8160
CAAGAGAAGC CCTGCCCGTT TTTCCAAAAC AGCCATAATC AAGCTAGCAC	GGTTAAAATC	8220

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AAGTCCTGTC GTAGTACGCT TGGCATTCC AAACATGGTC GGTGTTACCA AAGCCTGAAC	8280
CTCCGCCAAA ATCGGACGCG TCCCTTCAT GGTTACAACG ATGGAGGAAC CAGTCGCCCC	8340
ATCCAAACGC TCTTCTAGGA AAACTTGACT CGGATTGAGT ACCTCAACCA AGCCGCCCCA	8400
CTGCATCTCA AAAATCCCAA TCTCATTAGT GGAACCAAAA CGATTTTGAG CCGCTCTCAA	8460
AATACGAAAG GTGTGGTGAC GCTCCCTTC AAAGTAAAGC ACCGTATCCA CCATATGCTC	8520
CAACATACGA GGCCCAGCCA AGGTTCTTC TTTGGTCACA TGACCTACGA TAAAGATGGC	8580
AATGTTATTG GTCTTGGCCA ACTGCATGAG TTCAGCGGTC ACTTCACGCA CCTGAGAAC	8640
AGACCCCTGC ACCCTGAAA TCTCAGGAGA CATGATGGTC TGGATGGAAT CAATAATGAG	8700
AAAGTCTGGC TGGATACGCT CCACCTCTGC ACGAACACTC TGCACTATTGG TCTCTGCATA	8760
GAGATAAAAC TCACTATCAA TATCACCTAA GCGCTCTGCA CGTAGTTAA TCTGCTGGC	8820
AGACTCCTCC CCACTGACAT AGAGAACTGT CCCCACCTGG GACAACTGGG TTGAGACTTG	8880
TAGGAGAAGA GTTGATTTC CAATCCCAGG ATCCCCACCG ATAAGGACGA GACTTCCTGG	8940
TACCACTCCG CCTCCAAGCA CACGGTTGAA TTCTCCATC TCCGTTGGG TTGATTGAC	9000
ATTGATGGAA GTCACCTCAG CTAGTTCAT GGGCTTGGTT TTCTCACCTG TCAAGGACAC	9060
ACGCGCATTC TTAACCTCGG CAACCTCAAC CTCTTCCACA AAAGAACGACC AAGACCCACA	9120
GTTGGGGCAA CCTCCCAGAT ATTAGGGGA ATTATACCCA CAATTTGAC ATACAAATGT	9180
CGCTTTTTTC TTTGCGATGA CAAACCTCTT TCTATATCTC TAACTCACAC TCAATCACTT	9240
GGCAAAAATC AATCTCTCA TTTGGCACAA ACTGGCGCAT GAGCATTGCA TGAGCAACAA	9300
CTACCAACAGT CTGATGTTCT CGATACTTAG ACATACATTC TAGAAACCGA GACTTCATT	9360
CCGTAGCTGT CTCATATTGA ATAGGACTAT TAGGAAGCAA CTCCCCCTTG TTTTCTAAAA	9420
ACAGTCTTCT AGTGTGTTCA AAGTTTCTA TTCTGTTTT ATAGACCTGC CATTGATGTA	9480
ATAAAAGGCTC TACTCTTAAA GGAAGACCG TAGCACAGAC CACATACGAA GCCGTTCTA	9540
AAGCTCTTGT GACTGCAGAA GATACTGATTA TTTCAGCTGA CGAGAGTAA GGATTTTGC	9600
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CTATATAAGA ACGCTCCCT AACTCACGGT AATCTGGCTC CCCATGACGT ACAAAAGATAA	9720
TCTTCATTCT AGTGCCTGT CGATCCAAT CCACCAAGTTC GAACGCCATC AGCTGCATCT	9780
CCATCTGCAA TTAAGAAAAGT AGCAAAAACA GCCTGGACAA TACGCTCCCC AACTTCAAGA	9840
ACAACCTCTT GGTCTGTGAT ATTCTTCATC TGCGAAAAA TATGCCCTTC ATTTCCAGGA	9900
TTTCCATAAT AATCCCCATC AATGACTCCA ACTGAGTTAA TTAAAACCAA GCCCTTCTTA	9960

608	
CGAGGATTG AAGAACGATC ATAGAGGTAG AGAACCTCG AGAACCTCGAT ATAAGCCTTA	10020
AACCCCTGTCG GAACCAAGAC AATCTCTCCT GGCAGCAACAA CTGTACGAC AGCAACCTTT	10080
AAGTCGTAAC CAGTCGCATG CGCTGTCTCA CGCTTGGGCA ATAAATTTTC ATCTGTAAAAA	10140
CTCGAAACCA ATTCAAAACC ACCGAATTTTC ATAATTTCTCT TTTTCTATT ATCATTATT	10200
CTAGATTATT CTATACTTAT TTA	10223

(2) INFORMATION FOR SEQ ID NO: 74:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 16535 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 74:

TGGTTCTGTC CTTATCGCG CCTTGTCCTTG CTTGCCATGG CTACACCAAC TATCTCATCC	60
GACGAAACTA CACCAACCAC TAACGAACCC ACAACAGAA ATACAACCAC CCTTGCCCAA	120
CCTCTTACTG ATACAGCAGC TGCTCTGGT AAGAACGAAA GTGATATTTTC TTCACCTGGA	180
AATGCAAACG CTTCCCTAGA GAAAACAGAA GAAAACCTG CTGCAAGCCC AGCCGATCCA	240
GCACCACAAA CTGGACAAGA TCGTTCAAGT GAGCCAACTA CTTCTACTAG TCCAGTAACA	300
ACTGAAACATA AGGCAGAAGA GCCCATCGAA GATAACTACT TCCGTATCCA TGTCAAAAAA	360
CTTCCCTGAAG AAAACAAGGA TGCTCAAGGA CTATGGACTT GGGACGATGT TGAAAAACCA	420
TCTGAAAACCT GGCAAACGG AGCTTTGTCC TCAAGGATG CCAAGAAAGA TGACTACGGC	480
TATTACCTAG ATGTCAAATT AAAGGGAGAA CAAGCCAAGA AAATTAGCTT CCTCATCAAC	540
AATACAGCTG GAAAAAATCT AACCGGCGAT AAATCTGTAG AAAAACTACT TCCAAAAATG	600
AACGAAGCTT GGTTAGACCA AGATTACAAG GTTTCTCTT ACGAGCCACA GCCTGCAGGA	660
ACTGTTCGCG TCAACTACTA CCCCACAGAT GGCAACTATG ACAAGAAATC TCTCTGGTAC	720
TGGGGAGATG TGAAAAATCC AAGTAGCGCT CAATGGCCTG ACGGAACAGA CTTTACGGCT	780
ACAGGCAAAT ATGGCCGCTA TATGACATT CCTCTTAATG AAGCCGCAAG AGAATTGGAA	840
TTTTTATTAC TAGATGAGAG CAAACAAGGA GACGACGTGA AAATCCGTAA AGAAAATTAT	900
AAGTTACAG ATTTGAAAAA TCATAGCCAA ATTTCTCAA AAGACGATGA TGAATCGATT	960
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TCTAGCATTG AAAGTAGCTT TTCAACACTT GTCGGTGCTA AAAAGAAGA TATCCTCAA	1080
CACTCCAACA TCACTAATCA CCTAGGAAAC AAGGTAACCA TTACCGATGT TGCAATCGAT	1140

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GTTAGCTACA ATTCCGACCA ATTCACTACC AAAACAAGCT GGCGCCTGAA AGATGAGACA	1260
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CGTGAAGACG CCGTTATCTA CGAAGCTCAT GTGCGTGATT TCACTTCAGA TCCTGCCATT	1740
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TTGGAACCAA ACTACTACCA CTTTATGGAT GCCGATGGCA CACCTCGAAC TAGCTTTGGT	2160
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GCCGCTTCTA TCGAAGAAGC TTACAAGGCT GCACGCGCCC TCAATCCAA CCTCATCATG	2340
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TATCGGTAAT AACATCATCG ATAACCCAT TCCCCATTAT CTGGGCAAAA TCGCGCACGC	14340
GCTTGAGGAG ACGATTGGCA ATACGAGGGG TTCCACGACT ACGTAGGGCC AACTCAGATG	14400
CTGCCTCATG GGTGATTTCC ATCTCAAAA TATCTGCCGT CCGCTCGACA ATTTCTGTCA	14460
AGTCAGCATG AGCATAATAC TCCATATGAC CTGTAATCCC AAAACGTGCC CGTAGTGGAT	14520
TTGAGAGCAT ACCAGCCCCGA GTCGTCGCAC CAATCAAGGT AAAAGGAGGC AACTCCAAT	14580
GAACACTGCG ACTGCCCTCA CCAGCCCCAA TCATAATATC GATGTAGAAG TCCTCCATGG	14640
CACTATAAAG CACTCTTCC ACTGACATGG GTAAGCGATG AATCTCGTCA ATAAAGAGGA	14700
CATCTCCAGG CTCTAAATCA TTCAAAATCG CTACCAAATC ACCCGCTTT TCGATAACAG	14760
GACCAAGACGT TTGCTTGAGA TTGACTCCCA GTTCATTGGC ATGACAAAA GCCATGGTTG	14820
TTTTCCAAG CCCTGGAGGG CCAAATAAGA GCACATGATC CAGCGCTTC A TCCCGCATT	14880
TAGCGGCTTC GATAAGATC TGAAGTTGAT CCTTAACCTT ATCCTGACCA ATATATTAC	14940
GTAAATACTG AGGACGGAGC GTGGTCTA CTAACCTCTC ATCACCCATC ATCTCATTAT	15000
CTAAAATTCT ACTCATGGCT CTATTATATC AAAAAAAACA AGCCACAAAC AAAAAAGCCA	15060
CCTGATTGGG TGACTCCTAA GTTGTGACT TATGTGGTAT AATATTATAC GGCACTTCTA	15120
CACCGCCTAC GAAAGGAGGT GAGATAGCCC ATGATGAAAT TAGTACTCAA AACTATTATC	15180
GGACCAATTG TGGTCGGTGT CGTTCTTCGT ATAGTCGATA AATGGCTAAA CAAGGACAAA	15240
TAGTGTCAAA AAAGACCTCA AGCTTATTG GTCGTGAGCT TGGGTCTTT TCTAGCCTAT	15300

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GATATAGAAC TAGTACTCAA TTCCCTTTTA TTATCCCATA GTTCACGAAT TTTGTCAAAA	15360
CTTTACATTT CTCTCAACCG CTGTACGACA AGACGGTTAA GATTAAGAGA ACGTTAGGGA	15420
TTCTATCAAT TTCATAGAAA TTTTGATTTC GTAAACGAAG AGACAATCTT ACATGTCACT	15480
TCTCATTAA TACGCCACTA CTAGACAAGC AAAATCATTA TTACAGTAGT TCCAGTCCTT	15540
CAATTAACAG TCACTTACAA TCAAATTGAG TTTGAACTAG CTGAAGGCAC CACAGACCTA	15600
TTTCTTAGTC ATATTCGCTA AAAAAATCCC CGCCAAAATC TCAAAAAGTC CCCGCCAATT	15660
CCCCGACCAA AATCCGAAAA ATACCGAAAA ATATCGAAAA ATTATTTTTA GAATAGTCCC	15720
AAAAATCCTG AAATAGAGCT AAAAAACTCC ACCTGATTG GTGGAGTTAA GGGAGATTAT	15780
TATGAAAAAG AAAAGTTAG GATTTTATTA AATAAGTTA GGAGGTCTTT ATTTAATAAC	15840
TACATGATAC AAGACGAAAC TTAAAACTAG CTTAACTTTT CTAAAATTTT ACTATTTGC	15900
AAAAAAATTC TATCACCAGC ACCTCACCAA TCGAGTAGGG GATAATCTCT AGCCCCTCTC	15960
ACACCACCGT ACGTGCCGTT TGGCATACGG CGGTTCAACT AACTTTAAC GCATGTCGTT	16020
CAAGGTAATA ATCCAAACAC GAAACCAGTC CACGTTTTTC CAGGACTGGT TTTGATATAG	16080
CACGTTTAAG TACCGACTTC TGAGCTACTA ATTGATAATG GTGCCCCAG CCAGATACCT	16140
TATCTGCTAT CCATTTAGGA ACTCCTAACT TAAGCAATCC CCATAATCGT CTCGATTTCT	16200
TCTTCCATG CTTCCAGATA ATCACTCGTA GGGGAGTAGC CAAGGCTCA TCTATGCTGG	16260
CGACTATACT TTTCATATTT CCCAATGAGC AATAGTTTAT CCATCCTCGA ATAGACAAAT	16320
TCAGTTGCTC AATACTGCTT GTTAGGTCTA TACTCCATT CCTCTGTGTT AGTTTCTTCA	16380
ATTTAAACTT AAATCTCCGA ACACTATCTT GATGTGGACG GCTTTCCAA CCATCTGATA	16440
ATTTCCAGAA CCCAAAACCT AGATATTCA ACTCTCTTGG TCATGTTAC TTTCAAACCT	16500
AGCCGTTCT CAATAAACGA CTGACTGAAT ACATC	16535

(2) INFORMATION FOR SEQ ID NO: 75:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 8136 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 75:

CCAGAGCGTT GCGTCCGAAA GTCTATCCAG ACACGGCTCT TTAAAAACAA AAGGAGAAAT	60
GATGCATACT TATTTGCAAA AGAAAATTGA AAATATCAA ACAACCCCTAG GTGAAATGTC	120

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AGGTGGTTAC CGTCGATGG TTGCGGCTAT GGCTGATTAA GGATTTCAAG GAACTATGAA	180
GGCTATCTGG GATGACCTCT TTGCCCATCG TAGTTTGCC CAGTGGATT ATTGCTGGT	240
TTTAGGAAGT TTTCCCTCTGG CTGGAGTT GGTTTACGAA CATCGTATTG TTGACTGGAT	300
TGGGATGATT TGTAGCTTGA CAGGGATTAT CTGTGTAATC TTTGTATCGG AAGGTCGAGC	360
AAGTAATTAT CTTTTTGGCT TGATTAACTC TGTTATTAC CTTATTTGG CCCTACAGAA	420
AGGCTTTAT GGTGAGGTGC TGACGACACT TTACTTCACA GTCATGCAGC CAATTGGACT	480
TCTAGTTGG ATTATCAGG CACAGTTAA GAAGGAAAAG CAGGAGTTTG TCGCGCGTAA	540
ACTGGACGGC AAGGGCTGGA CAAAGTATCT TTCCATTAGT GTGCTTTGGT GGTTGGCCTT	600
TGGCTTCATT TATCAGTCTA TTGGTGCCTAA TCGTCCCTAT CGTATTCAA TCACAGATGC	660
AACCAATGGG GTAGGGCAAAC CCCTCATGAC AGCTGTTAC CGTGAACAGT GGATATTCTG	720
GGCGGCTACC AATGTCTTT CAATCTATCT CTGGTGGGAA GAAAGCCTGC AAATTCAAGG	780
GAAATATCTA ATTTATCTA TTAACAGTCT AGTTGGTTGG TATCAATGGA GCAAGGCAGC	840
TAAGCAGAAT ACTGATTTAC TTAACTAGGA AAAGATGTTT GAAAGTGCCTG TTTTGAGATT	900
TCGATTAAGG CAGATATACT TGATAATCAA GGATTTATAG TATGAAAAAG AGGATCGCGC	960
GGTCCTCTTT TGTTGTTGAA AAGATAAAA ACTCAGTAAC CTAGAAATAA GACAACGTGAA	1020
GCTTTACTCT ATATTCAATT TTTAGGAATG AGAAGGTCTA GATAAAATTG GACAACCTCC	1080
TGGTCTGTGA AATCTTGACC TTTTTGAGC CACCAGGTCA ATGTCTCGAT AAAGTTGGAC	1140
ATGACCAAGT GTTGGAGGTA AGAACTAGGC AGATTAGGGT GGGCTTCTTT TAAATTATCA	1200
GCTAGCACGG AATAGACATG GTGTTCTAGC TCTTTATGGA GTTGACGGAG GAAGTAGTCA	1260
TTTTTGAAA ATAGCAGACT GGTGATATGG TCTTGTTTT TATGAAAATG GAGAAAGAGG	1320
TGGGCGAGGT AGTCCTCGGT TGAAATGGCT TGCTCTCTTT CAAAAAGATG ATGGAAGAGG	1380
TAGCGGCAGA GCTGGTCCAG AAGAACGCTCC TTACTCTCAT AGTGCACAGTA AAAGGTGGAT	1440
CGTCCCACAT CTGGCAGATC AATGATATCC TGAACAGTAG TGGCCTCGTA GCCCTTAGCA	1500
TTCAAAAGTT GTATAAAAGC TTGATAGATG GCTTTTTGG TTTTGCTGAT ACAGCGGTCA	1560
ATGTTAGTCA TATGGACACT TAAGGCAAAT TGTCAGAAC TGAATAAAGC TGACGTTTG	1620
CTTCTATCCT TTCTTGAGT TTTAGTGGAT AATGATAATG ACAAGGTGT TCATAAAATCT	1680
ATTATAACAA AGGAATGAGA AATATGAAGG CAAATATGC TGTTGGGTG GCTTTTTCT	1740
TAAATTGAC TTATGCCATT GTTGAGTTA TTGAGGTGG AGTATTGTT TCTAGCGCTG	1800
TTCTTGCTGA CTCTGTGCAT GACTTGGGAG ATGCGATTGC AATTGGAATA TCAGCTTTTC	1860
TAGAAACAAAT CTCCAATCGT GAAGAAGACA ATCAGTACAC CTTGGCTAT AAGCGGTTA	1920

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GCCTGCTAGG AGCCTGGTA ACAGCTGTGA TTCTCGTAAC GGGCTCTGTT CTAGTCATT	1980
TGGAAAATGT CACGAAGATT TTGCATCCGC AACCAAGTCAA TGATGAGGGG ATTCTCTGGT	2040
TAGGAATTAT TCAGGATTACT ATCAATCTGT TAGCGAGTCT GGTGGTTGGT AAGGGAAAGA	2100
CAAAGAATGA GTCTATTCTG AGTCTGCATT TTCTGGAAGA TACGCTAGGG TGGGTAGCTG	2160
TTATCCTGAT GGCGATTGTT CTTCGATTAA CGGACTGGTA TATCCTAGAT CCTCTTTGTT	2220
CCCTTGTCAT TTCTTTCTT ATTCTTCAA AAGCCCTTCC ACgtttttgg tctacactca	2280
AGATTTCTT GGATGCTGTG CCAGAAGGTC TTGATATCAA GCAAGTAAAG AGTGGCCTGG	2340
AGCGATTGGA CAATGTGCC AGCCTTAATC AGCTTAATCT CTGGACTATG GATGCTTTGG	2400
AAAAAAATGC CATTGTCAT GTTTGTCTAA AAGAAATGGA ACATATGGAA ACTTGAAAG	2460
AGTCTATTCTG AATTTCCCTA AAAGATTGTG GTTTCAAAA TATTACCATT GAAATTGATG	2520
CTGACCTAGA AACTCACCAA ACCCATAAGC GAAAGGTGTG TGACTTGGAA CGGAGTTATG	2580
AGCATCAACA TTAGAAAAAA GTGAAAAATA CTTGGGTACT ATCTTATTG GAATAGAGTA	2640
ATTTCTTTAT TATTTAAATA TTTCAAAAT TGTAAGAGA AGAGCATTGT ATAAACTCCA	2700
GATATATGAT TGTTAATGAT AAAAATTTT CGATTAGATA CAAATGCTT GACTTGGAGT	2760
CAACTCAAAG TTATATAATA AGATAAGTGA GTTAAATAG CGTGAATTCA GTGAATGAAA	2820
TGAGAGGAGG TTAGCGTGTG AATATTAAT CTGCCAGTGA TTGTTGGGA ATTTCAAGCGG	2880
ATACGATTGCG TTTTATGAA CGGTTGGTC TTGTGCCACC GATTAATCGT ACTGCTACTG	2940
GGATTCGTGA TTTTCAAGAT CAGGATATCG AAGCGCTGGA ATTTATTAAG TGTTTCGTT	3000
CGGCGGGTGT CTCTGTAGAT AGTTTAGTTG ACTATATGTC GCTCTACCAA AAGGGAGATG	3060
AAACGAGAGA GGAGAGGCTT GGTATTTAG AAGAGGAAAA GCAAAATTA GAGGAGCGCT	3120
TGTCTCAGCT ACAGACAGCT TAAATCGTT TAAATCTAA ATTAAACTT TATAAGGAAG	3180
GAAAATTTA AATGAATCA GCAGTATATA CAAAGGCAGG TCAGGTTGGA CTTGCTAGCA	3240
TTGAACGTCC GCAAATAATA GAAGCGGATG ATGTGATTAT TCGTGTGGTT CGTGCCTGCG	3300
TTTGTGGTTC AGATTTATGG AGGTACCGTA ATCCAGAAC GAAAGCTGGA CACAAAATA	3360
GTGGACACGA AGCGATTGGG ATTGTTGAAG AAGCTGGGA AGCCATTACG ACGGTGAAAG	3420
CAGGTGATTT TGTGATTGTC CCTTTACAC ATGGATGTGG TGAGTGTGAT GCCTGTCTG	3480
CTGGATTGAA CGGTTCTTGC GACAATCATA TTGCAATAA TTTGGGGGGT GATTTTCAGG	3540
CAGAATATAT TCGCTTCCAC TATGCAAACG GGGCGCTGGT TAAATCCCT GGTCAACCTT	3600
CTGACTATAC AGAAGGGATG CTCAAGTCCC TTTGACTCT TGCAGATGTC ATGCCGACAG	3660

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GCTATCATGC GGCGCGTGT GCAAATGTT AAAAGGGGA CAAGGTTGTT GTTATCGGTG	3720
ATGGGGCTGT TGGTCATGT GCTGTCATCG CGGCTAAGAT GCGTGGAGCA TCACAAATTA	3780
TCCTTATGAG CCGTCATGAA GACCGTCAAA AGATGGCTAT GGAGTCAGGT GCGACAgcTG	3840
TTGTTGCAGA ACGTGGTCAA GAAGGAATTAA CCAAGGTGCG TGAAATCCTC GGTGGAGGAG	3900
CAGATCCAGC ACTTGAATGT GTTGGTACGG AGGCTGCTAT AGAACAGGCG CTAGGTGTT	3960
TTCATAATGG AGGGCGTAGT GGCTTGTTAG GAGTCCCACA CTATAATAAT CGTGCTTTG	4020
GTTCGACATT TATGAAAAT ATCTCTGTAG CAGGTGGGC AGCTTCTGCT ACAACATACG	4080
ATAAGCAATT TTTACTAAAA GCCGTCCTTG ATGGTGATAT CAATCCAGGT CGCGTCTTTA	4140
CTTCAACTTA TAAACTGGAA GATATCGACC AAGCCTATAA AGATATGGAT GAACGTAAGA	4200
CAATTAAGTC TATGATTGTA ATCGAATAAA AAACGAATAG GAGTTTTAGA ACTCTATTG	4260
TTTTTTATGT TATCCTATTTC TTGATTAGG GTACTTTCTC TTAATGTCAG TCTGGTTCCC	4320
AGCATGGTCA GGCTAGGGAT TTTCCGACCG TGGAGGACTT CCTTGTAAAG AATATCCATA	4380
CCTGCTCGGC CCATTTCTTC AGTATAAACT GTAATACTAG AGAGGGGAGG ATAGACCTGT	4440
TTGGTCAGAC TAGTGTGTT AAAGGAAATG AGGCTGACGC GATCTGGCAG GCTGATTCCA	4500
GCTTCTTGGA GGGCACGGAG GGCACCGATA GCTAAACTAT CGCTGGCTGC GAAAAATGCT	4560
GGCGGAAGTT GGTCTCCCAA GCTCTGAATG GCCTCCTTCA TTAAGTCATA GCCAGACTGG	4620
GCAGTAAATC TTCCTTGAAA GACCAGTTCA TCATGATAGA TTCCCCTCGC TTGACTATAG	4680
TTTTTGAAAGT TTTCTAGACG CTTGTCCTGA ATGATTCTT CTTGGTCTGT TGTTTCTTC	4740
AGGCCCTGTTA GAATCCCGAT ACGGTCCATT CCTTGACTGA GGAAATAATC GACAACCTGT	4800
TTCATAGCAG TGTAAAATC CGTGATAATA CAGGTATGTC CCAGGGAAAG TGTATCGCTG	4860
TCTAGAAATA CAAGAGGCTT TTGGTATTCT TCAAAGGCAG AAATCTGAGC TCGACTAAC	4920
TTTCCGATGC AGAGAAATCCC AATCACTTCC TCGCTTAGGG TAAAAGGTG GTCATTAAAA	4980
TAGGCCAAGA TATCATAGTC CAACTCTTGG GCTCTTTTT CTATTCCTAG GCGAATCTGG	5040
TAGTAGTAGA GGTGGTCCAG CTCCCCTTGT TCGCTGACCC ATTGGATAAT GGCAATCTTT	5100
TGCTTGGGTT TGTGGGACTC GCCTGTCTTG AGGTGCTTGG TGTAGCCCAG CTCTTCAGCA	5160
ACGGTTAAA TACGGTGTCT GGTTTCTCT GTAAACAGATA GGCTCTGGTC GCGGTTGAGG	5220
ACGCGGGATA CGGTGCGCAT AGAGACAGAG GCTAGCTGTG CAATGTCTTT TAAGGTAGCC	5280
ATAAATCCTC CTTGATTAGG TTAGTATATC ATGTTTTCT TCTTTTACT GATATTTAC	5340
AAAAATTITA GTAAAAGGA TTGACCTTGG AAAATCCTT GGATATAATA GAAAGAAAAC	5400
GATTACACGT TAAGATGGCT TAACGGACAG TCAAAGGAGA ATTCAATATGG CACAAACATCT	5460

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TACTACTGAA	GCCCTTCGCA	AAGACTTTCT	TGCTGTTTTT	GGTCAAGAAC	CAGATCAAAC	5520
CTTCTTTCA	CCAGGCCGCA	TTAATTGAT	TGGTGAACAC	ACAGACTACA	ACGGTGGGCA	5580
CGTTTTCTC	GCTGCTATT	CCTTGGGAAC	TTACGGTGCA	GCTCGTAAGC	GTGACGACCA	5640
AGTCTTGCCT	TTCTACTCAG	CTAACCTTG	GGACAAGGGC	ATTATCGAAG	TGCCTCTCGC	5700
TGACCTCAAG	TTGAAAAAG	AGCACAACTG	GACCAATTAT	CCAAAAGGTG	TCCCTCATTT	5760
CTTGCAAGAA	GCTGGGCACG	TGATTGACAA	AGGTTTGAT	TTTTATGTTT	ATGGAAATAT	5820
TCCAATGGT	GCTGGCTTGT	CTTCTTCTGC	ATCCTTGGAA	CTCTTGACAG	GAGTCGTGGC	5880
TGAGCATCTC	TTTGATTTAA	AATTAGAGCG	TCTCGATTTG	GTAAAGATCG	GCAAACAAAC	5940
AGAAAACAAAC	TTTATCGGAG	TAAACTCTGG	CATTATGGAC	CAGTTGCTA	TTGGTATGGG	6000
GGCAGACCAA	CGTGCTATT	ACCTAGATA	TAATACTTTA	GAATACGACT	TGGTGCCACT	6060
TGATTGAAAG	GACAATGTCG	TTGTTATCAT	GAACACCAAC	AAACGCCGTG	AATTGGCGGA	6120
CTCTAAATAC	AATGAACGTC	GTGCTGAGTG	TGAAAAGCA	GTGGAAGAAT	TGCAAGTTTC	6180
CTTGGATATT	CAGACTCTGG	GTGAATTGGA	CGAGTGGGCC	GTTGACCAAT	ATAGCTATCT	6240
GATTAAAGAT	GAAAATCGTT	TGAAACGTGC	TCGCCATGCT	GTGCTTGAAA	ACCAACGTAC	6300
CCTCAAAGCT	CAAGTAGCAC	TCCAAGCAGG	AGATTGGAA	ACATTTGGAC	GCTTGATGAA	6360
TGCGTCACAC	GTTCCTCTGG	AGCATGATTA	TGAAGTAAC	GGTTTGGAAAT	TGGATACCT	6420
TGTTCACACA	GCTGGGCAC	AAGAAGGGAGT	TCTCGGTGCT	CGTATGACAG	GGGCTGGTTT	6480
TGGTGGCTGT	GccATTGCCT	TGGTTCAAAA	AGATACTGTT	GAGGCCTTTA	AGGAAGCTGT	6540
AGGCAACAC	TACGAGGAAG	TAGTTGGATA	CGCTCCAAGC	TTCTATATCG	CTGAAGTTGC	6600
AGGTGGCACT	CGCGTCCCTG	ACTAGTCAA	AGGAGGCTCT	ATAGTGACCT	TAGTAAATAA	6660
ATTTGTAACA	CATGTCATT	CTGAAAGCTC	ATTGAGGAA	ATGGATCGAA	TCTATCTGAC	6720
CAATCGTGT	TTGGCACGAG	TGGGAGAAGG	TGTTTGGAA	GTTGAGACCA	ATCTGGATAA	6780
ATTGATTGAC	CTCAAGGACC	AGCTGGTTGA	AGAAGCCGTT	CGATTAGAGA	CGATTGAGGA	6840
TAGTCAGACT	GCGCGTAAA	TCCTTGGTGC	TGAAGTGATG	GATTGGTGA	CTCCTTGTCC	6900
AAGTCAGGTC	AATCGTGATT	TTTGGGCAAC	CTACGCCAC	TCTCCAGAAC	AAGCGATAGA	6960
GGATTTTAC	CAACTCAGTC	AGAAAAATGA	CTACATCAA	CTCAAGGCCA	TTGCTAGAAA	7020
TATCGCTTAT	CGTGGTCCAT	CTGACTACGG	AGAACTTGAA	ATTACCATCA	ATCTCTCTAA	7080
GCCTGAAAAA	GATCCCAAAG	AGATTGTGGC	AGCCAAGTTG	GTGCAAGCTA	GTAATTATCC	7140
TCAGTGTCA	CTTTGCTAG	AGAATGAGGG	CTACCATGGT	CGAGTTAAC	ACCCAGCTCG	7200

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TAGCAATCAC CGTATTATCC GTTTGAAAT GGTTGGTCAG GAATGGGTT TCCAGTATT	7260
GCCCTATGCT TACTTTAATG AGCATTGTAT CTTTTAGAT GGCCAGCATC GTCCCATGGC	7320
CATTAGTCGT CAGAGTTTG AACGCTCTGTT GGCTATCGTA GACCAGTTTC CAGGATATT	7380
TGCTGGATCT AATGCCGACC TGCCGATTGT GGGGGCTCT ATTCTAACTC ATGATCATTA	7440
TCAGGGAGGC CGTCACGTAT TTCCTATGGA ATTGGCTCCC TTGCAAAGG CCTTCCGATT	7500
TGCTGGTTT GAGCAGGTCA AGGCTGGAAT TGTCAAGTGG CCCATGTCCTG TCCTACGTT	7560
GACTTCGGAT TCCAAGAGG ATTTGATCAA TTGGCTGAT AAGATTTCG AGGAATGGCG	7620
CCAGTATTCA GATCCTGCAG TGCAAGATTT GGCAAGACA GACAGGACAC CGCATCACAC	7680
TATCACACCC ATTGCCCGCA AACCGGATGG ACAGTTGAG TTGGACTTGG TCTTGCGAGA	7740
CAATCAGACT TCAGCAGAGT ATCCTGATGG TATCTATCAT CCCCCACAAGG ATGTCCAACA	7800
TATCAAGAAG GAAAATATCG GCTTGATTGA GGTCAATGGC TTGGCAATCT TGCCACCACG	7860
TCTGAAAGAA GAAGTGGAGC AAGTCGCTAG CTATCTTGTA GGAGAAGCTG TTACAGTTGC	7920
CGATTATCAT CAGGAGTGGG CAGACCAACT CAAATCCCA CATCCAGACT AACGGATAAA	7980
GAAAAAGCCC TTGCAATCGT CAAGGACTCT GTGGGTGCTA TCTTGCAGCG TGACTTGAG	8040
GATGCAGGAG TCTACAAGCA GACAGAACAA GGGCAGACAG CCTTTATGCG CTTTGTGGAA	8100
CAGGTGGAA TTTTACTAGA CTAGGAGCTT TCTCGG	8136

(2) INFORMATION FOR SEQ ID NO: 76:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 10011 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 76:

CCCATAGTGA AGAGTGGCCA TAAGAAGGTC TTCTAGGCTT AATTTAGGTT TTGTCAC	60
TTTTGCGTGT TTAAGTTGAT AAGCTGTTT TAACACAGCT AACATCTCT TCAAAAGTCG	120
TGCGCTGAAC ACCAACAAAGA CATTAAATC GTGTATCAGT TAGTTGTTA CTTGCTTCAT	180
CATTCACTAGA ACTACTATAC CATGTTTGTT TTCCGAGGAA GTCTAATATT GTCAAATACT	240
GGAACGCTCA TTGCTGGAT ACGGAATAAG ATTGGCCAG CTTCGATAAC TGGGATACCT	300
GGTTCAAAAC CAAGGTCTGT TGCAAGCGATT GGTGTAAAGA TATCGTAACC TTTCTATAAGG	360
TCTTCGTTA CATCTTCAC CATAACTGCA TCACAGTGA CATCGTAACC ACGGTTGAA	420
AGTTCTCTT CTAGAGCACT TTTAATTGAG TGACTTGACT TAACACCTGC ACCGCAGGCA	480

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GCAAGAATTT	TAATCATTG	GATTCCTCC	GATTTTATTT	TTTAATAGAC	AAGATTAAGC	540
GGTTGCTTCA	GCAATGTAAG	CATAAAGGGC	TTCTGGTCA	GAAATTTTG	ATAGGTCTTC	600
AAGATGACCA	TTTCCTGTGA	AGAAGTCCAT	TAACTGAGCA	AGAATGTTCG	TTTGACTTGA	660
ACTTGAATTA	TTGATGATAA	AGAAGAGCAA	GGATACTTCT	ACTTCCTTAC	CTGGCGCAAT	720
CATATTATGG	AAAGTCACCG	TTTCTCTAA	TCGAACAAACC	ACCACTTTCT	CAGCTAGATT	780
ATGAACAATA	TCTGTGTGAG	GAATCATTAC	ATTTGCAAGT	CCTTCCTAG	AAATTCCATA	840
TATAAACCCAG	TTGGAATGA	CTTTTCACGC	GTGATCAAGG	CTTCACGATA	AGTTGGAGTG	900
ACAATTCTC	GTTCTTCAA	CAAGCTTGCT	ACCTGATCAA	AAAGTTATTC	TTGATTATCC	960
GCTTCTAACG	AAAACACAAG	GTTTTGTCA	AAGAAATAAT	CTAATACCAT	AAGGTTTTCC	1020
CTTCTTCCA	TTAACTTTAT	GCTATAAGTA	TAACACTATA	TGAAATCGTT	GTTAATTACT	1080
TTCTATTCTT	TTTTGCTCT	TTTTTTATAT	TTTGTTTTG	TTTATAGTTT	GTTATATAAA	1140
AATAAACACA	CAAACAAATA	CTCCAAGCAT	TTTTCTGTT	TAATACTCAA	TGAAAATCAA	1200
AGAGCAAACT	AGGAAGCTAG	CCGCAGTTGT	TCAAAACACA	GTTTTGAGGT	TGTAGATGAA	1260
ACTGACGAAG	TCACTCAAAA	CATGGTTTG	AGGTTGTAGA	TGAAACTGAC	GAAGCAACAg	1320
CCATACATAC	GGTAAGGCGA	CGCTGACGTG	GTGAGAG	ATTTTCAAG	AGTATAAAAA	1380
CTAAAAAAGC	AGACCATCTA	AGCCTGCTTT	ACTATTGATT	CTTATATAAA	TTTCTGTGA	1440
ACAAGGAAAG	GCATTTCTGA	TAACTTATTC	TTCATCCATA	CTCAAGACGC	TGAGGAAGGC	1500
TTCTTGGGA	ACTTCAACTG	ATCCGATGGA	TTTCATGCGT	TTCTTACCAAG	CTTTTTGTTT	1560
TTCAAGGAGT	TTACGCTTAC	GAGAACGTC	ACCACCAAA	CATTAGCAA	GTACGTTCTT	1620
ACGAAGGCC	TTGATATCG	TACGAGCGAC	AATCTTGTT	CCAATAGCCG	CTTGGATTGG	1680
AACTTCAAAT	TGTTGGCGAG	GGATGTTTT	CTTGAGTTTA	TCAACGATGA	GTTCCCCACG	1740
TTCCTAGGCA	AAGCCTTGT	GAACGATAAA	GCTGAGGGCA	TCCACCTTAT	CTCCATTGAG	1800
AAGAATATCC	ATTTTCACCA	GCTTAGATGG	GCGATATTCT	GACAATTCTG	AGTCAAAGCT	1860
TGCATAACCA	CGTCTCGAAG	ACTTAAGTTT	ATCAAAGAAC	TCAAAGACAA	TTTCAGCAAG	1920
AGGAATTG	TAGATAAACAT	TGACACGGTT	ATCATCAATA	TAGTCCATAG	TCACAAAGTC	1980
CCCACGCTTA	CGCTGAGCTA	GCTCCATTAC	TGCTCCGACG	AACTCCTGTG	GTACCATGAT	2040
TTGCGCCTTG	ACATAAGGCT	CTTCAATGGT	CGCAATCTTA	GTTGGGTCTG	GAAACTCAGA	2100
TGGGTTAGAC	ACATCCATAG	ACTCACCGTC	GGTCAAATTA	ACTTTGTAAA	TAACAGACGG	2160
AGCTGTCATG	ATGAGGTCAA	TATTGAACTC	ACGCTCTAAA	CGTTCCCTGGA	TAACATCCAT	2220

624	
ATGGAGAAAGT CCAAGAAAATC CACAACGGAA ACCAAATCCA AGTGCCTGAG ATGTTTCTGG	2280
TTCAAACTGA AGACTAGCAT CATTCAAGTTG CAATTTTTCA AGcGCTTCAC GCAGGTCATT	2340
GTACTTGTTC GATTCGATTC GGTAGAGACC CGCAAAGACC ATAGGATTCA TCTGCTTATA	2400
ACCATGTAAT GGTTCTGCCG CAGGATTGGT TGCCAAGGTA ACGGTATCAC CCACACGAGT	2460
ATCCTGAACC GTCTTGATAG ACGCCGCAAT GTAACCAACA TCACCAAGTCG CAAGGAAATC	2520
ACGACCAACC GCTTTGGTG TAAAAATACC GACTTCGGCC ACATCAAAGG TCTTACTATT	2580
GCTCATGAGC TGAATCTTAT CACCAAGTTT GACCACTCCG TCCATGACAC GCACATTGGAG	2640
GATAACCCCA CGGTAAGCAT CGTAAACAGA GTCGAAAATC AAGGCCTTAA GTGGCGCCGT	2700
CACATCACCC GTTGGTGCTG GTACTTTTC TACAATTTGC TCGAGGATTT CTTCAATCCC	2760
AATACCAGCC TTGGCAGAAG CCAAAACTGC TTCACGGCA TCCAAACCAA TCACATCTTC	2820
AATCTCTGTA CGCACCGCCT CGGGATCTGC AGCCGGCAGG TCAATTATTA TAATGATAGG	2880
CATGATTTC CAAATCATTAT CCAAGCCAG ATAAACGTTG GCAAGAGTTT GAGCCTCAAT	2940
TCCTTGAGCC GCATCGACCA CCAAAATAGC ACCCTCACAG GCAGCTAGCG AACGTGAAAC	3000
TTCATAGCTA AAGTCAACGT GCCCTGGTGT GTCAATCAAG TGGAAAATAT AAGTTCCCC	3060
ATCTTTGCA GTGTAATTCA ACTCGATGCC ATTCAACTTA ATAGTAATTG CACGTTCCCG	3120
CTCTAGCTCC ATGCTATCCA AAAGCTGGC CTGCATTCA CGACTTGAAA CCGTCTCTGT	3180
TTTTCCAAA ATGCGGTCTG CTAGAGTTGA TTTTCCGTGG TCAATATGGG CGATAATAGA	3240
GAAGTTACGG ATCTTCTCCT GTCGTTTTT CAATTCTTCT AAGTTCATGA TTCTCTTCCT	3300
TTCAGGGTAT CTATTATTAA TAAATTGTTT TTGATATTGTT GACAAGACCA TACCCGTCTA	3360
GGAGTACTAA TCTTCAGCGA CAAAGCCGTC ATTTTCGATA AAGTGGTGT CTGTCATTCC	3420
TTGGTCTGTA AAGACAATCC CGTGAAGGAC ACCACCAAA ACAGCTCCTC CATCCATTCC	3480
AATCTTGCCA TCTTCTGTAG TCCAAAGCTC AGATGTACCG CGTTCTTGCT GTAACAAACC	3540
ATAGACCGGT GTATGACCGA AGACAATGGT TTTTCCAGTA TGATTTTCAG CTCCGTGGAA	3600
TGGTTTTCTA AGCCATACTT TTTTATAATC TGTTGTTCA TGCCAGTCGT CCAAGGTCAA	3660
ATCAATAACCT GCGTGAACAA AGATATACTT GTCTGTCTCT ACTACAAATG GCATTTGACG	3720
AATGAATTG ACCAAGTCTG CCCGTTCAAG GgCAACCCGC TTGGCATCTT CTACTCCATC	3780
AACTGGTGCA TCCAAGGGAC GACCTAGGAT AGAGTTAATG GTTGTATCTC CACCATTGCG	3840
ACTATAATGG TCATAACTTT CTTCTGGTC ATCTAGCCAA GTCAAAAACA TATACTCGTG	3900
GTTCGGGAC AAACAGATAG CCCCTTGATT GTCCACCAAG TCCTTGACCA TTTCAAGAAC	3960
ACGGTGACTA TCCTCACCTC TGTCAATCAA ATCACCTAGA AAGAGCAACT GGGGCTGACC	4020

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ATCCCAGGTT TTGAGAAGGT CTTCCAGCAT CCCAGCTTT CGCGAACAT CTCCAATTAC	4080
ATAATAATCT GTCATCTTAT TTCTCCCTGT TTCTCAACAA TTCTCTTGCT TGCGTCAGGG	4140
CTGCTCTGTG CACATCATCA CCTGCCAACA TCTTGGCAAC TTCCCTCCACT CGCTCTTCGA	4200
CCGTCAAGAG ACGAACAGTC GAAACCGTTG AATGGTCATT ACTAATCTTC TCAATAAAGA	4260
ATTGATAATC TGCAATCGCA ATTACTTGTG GCAAATGGGA GATAGCCAAA ACCTGACCAC	4320
GCTGACCAAT TTTATGAATT TTCTGACCAA TAGCTTGAGC AACACGACCT GAAACTCCCC	4380
TATCCACCTC ATCAAAGACA ATGCTAGTCT TGCCTTCTTT ACGTGAAAAG GCAGACTTAA	4440
TGGCTAACAT GAGACGGAGAT AATTCCCCTC CAGAAGCAAC CTTAACCAAG GGTTAAAGT	4500
CTTCTCCAGG GTGGGTGAA ATATAAAACT CAACCATTTT ATTTCCCTCA CGACTGAATT	4560
TTCCCTTACT AAAACGAACC TGAAACTGGG CTTTTCCAT ATAAAGATCT TGCAGTTCTT	4620
GTAAATCTC AGCTTCCAGT TGCTGAGCCA AATTATGACG AGCAGAAGCA AGTTGACCTG	4680
CCAAATTGAC AAGATTGACT TCCAACCTCT TAAGCTCTGC TTCCATGTCC TCAGACGAAA	4740
GATTATTGCC TGTCAAGAGA TTGTATTCTT CCGTAATCTT GGCAAAATAA AGCAAAACAT	4800
CATCAACAGT CCCACCATAC TTACGAGTAA TAGTATGAAG GAGGTCCAAA CGATTCTCAA	4860
CCTGCATCG GCGATTGCCA TCAAAATCAA GGTCCTCAAT GATAGCTTCC AAACGTTTGC	4920
TAATGTCCTC TAAAACATAG TAGGTCTCG ACAGATACCT TGAATTTCA CGGTATTCA	4980
GATCATACTC TTCGACACTT TCCATGTCAT TCATAGCTGA ACGAACATTG GCCAGACTTG	5040
AAAAATCTTC ATTGTCACAC ATACTGTAGG CATTGGTCAG TGTATCCGCA ATATTTTTGT	5100
GGTTGAGGAG TTTATCTCGC TCTTGATTGA GAGCCAAGTC TTCTCCAGCC TGCAAGTTTG	5160
CTGCCTCAAT CTCTGCCATT TGAAATTCCA ACATTTCGAT ACGTGCCTTG TGTCCTGTT	5220
GGTTTTCTT GACTTCCAGA ACCTGCTTGC GCATTTCCG ATAGGCATCA AAACCTGTT	5280
GATAGGTTTC TTTCAAGTCC CAAAAAGCGG CATCACCAAA TTCATCCAAC ATCTGGATAT	5340
GCAGTTGGGG ACGCATTAAC TCCTCATGGT CATGCTGACC ATGAATATCT ACAAGATGTT	5400
GCCCAATAGC TCGAAAACA GACAGATTA CCATCTGACC ATTTACACCG CTGATACTAC	5460
GACCATTTCG CAAGATTC CGACGGATGA TAATTCATC ACCTAATTCT AACACCTTGCT	5520
CATCAAAAAT TTCTGTAAA AGACGACTAT TCTCAACTGA GAAAAGCCCC TCAATCTCG	5580
CCTTTGGTGC ACCATGACGA ATAACATCTG TCGTCGCACG AGCTCCCAAC ATCATATTCA	5640
TGGCCTCAAT GATAATCGAC TTCCCTGCAC CCCTTCACC AGTCAGGACA GTCATCCCC	5700
TTTCAAAATT GAGGGAAATA GCCTCAATAA TGCCAAAGTT TTTTATCGAA ATTTCAAGTA	5760

626	
ACATATAGAC CTACCAATT TTTACTTGT CAAAGATTC CTCTGCTAGA CTTCCACTTC	5820
TGGCAATGAC TAAAATCGAG CTATCATCAG TCAAACAGCT AAAAATCTTG TCTGCAAAG	5880
TCTCGATTAA CTGAGCTTT ACAAAAGCCG TATTTCTGG AATAACTTGG AGATTGATCA	5940
TCTTATCCAT CAATTCAAGCC GATTCGATAT TGTCTTCAGC CAGTTGCAGA CTTTTTACGA	6000
TTGATTTGG CAATTCTGAG ACATAGGTGT TGTCTCTCAA AGGAATTTG ACAATACCTA	6060
ACTCTTGAT ATCTCGGGAT ACCGTCGCCCT GAGTGGCAGT GATACTGCT TCTTTCAAAT	6120
GTTCTACAAT TTCTTCTTGC GTGCCGATTT GATAATCTGT CACCAATCTT CTAATTTTTT	6180
CAAGTCTCTC TTTTTTATTG ATTTTTAAAT TGACTATGCG CCCTCTCTAC TGCTTCTTTA	6240
ATCTCAGCAA GAATCTGATT GCTTGCTGAC TTTTCTTTT TCAAATACGC TAAAAATTCA	6300
ATATTTCCAT GTCCACCTTG GATGGGAGAA AAGTCCAAGC CAAGGACTGA AAAACCTACC	6360
TCTACTGCCA TAGCTGTTAC AGATTCAGG ACATCTGAT GAACCTTAGC ATCTCGAATA	6420
ATTCCATTTT TCCCACATCTG CTCACGTCCT GCCTCAAACG GAGGTTTGAC AAGTGCTACC	6480
ACCTGACCTT GATCAGCCAA GACACGGTGC AAGGCTGGCA AAATCAGACT AAGGGAAATG	6540
AAACTCACAT CAATACTGGC AAAGCTCGGC TCTGCTCGA AATCAGTCTT TTCAGCATAG	6600
CGGAAATTGA ACTGCTCCAT GCTGACAACG CGTGGGTCTT GGCGTAATTG CCAAGCCAAC	6660
TGATTGGTAC CAACATCGAC TGCAAAGACC AACCTGGCAC TATTCTGTAG CATGACATCG	6720
GTAAAACCTC CAGTAGAGGC CCCGATATCA ATCGTAGTCG CGCCATCCAC CGACAAATCA	6780
AAGACCTGCA AGGCCTTTTC CAGTTCAAA CCACCACGGC TGACATACTT GAGTTCTCC	6840
CCCTTGAGTT TTAATTCTGGT GTCATCTGGA ATTTCTCTC CTGGCTTGTC AAACCGTTCT	6900
CCATTAAGGA CTGCTACGAC TAGGCCAGCC ATCACACCTC GCTTGGCCTG CTCTCTCGTT	6960
TCAAACAAACC CCTGTTATA AGCTAGTACA TCCACTCTT CCTTAGCCAT TGATTCTCAA	7020
ACTTTCTACT ACACCTACAA TCGATTCTGT TTCAAAGGGA AGCTGCTGGG CAATTCTTC	7080
TAATTTTCA TTAGCTTGT CCAGGGTTTG GTTACAAAAG GCAATGGACT CTTCCAAGCC	7140
CAACAGGGCA GGATAGGTTG ATTTCTGC CTGCAGATCC TTTTGAGGTG TCTTGCCGAT	7200
TTCCCTAAAA CTAGCTGTCA CATCCAGTAC ATCATCTCTG ACTTGAAAAG CAAGTCCAAT	7260
CAATTCAACCC ACAGTTTCA GCTTCACCTG CATTTCAGGT GACAATTCAAG CTATAATAGC	7320
TGCCGCTTGG AAGGGATAGG CTAGTAACCTT CCCAGTCTTA TTGGCATGAA TAGTCTGAAG	7380
TTCTTCCAAA GACAAGTGCT GGTGTTCGCC CTCCATATCC AAAAATGCG CTGCTACCAT	7440
ACCCAGACTA CCTGAAGCAA GGGATAAGTT GGCAATCAAG TCCACCTTAA TCTGACTTGG	7500
CAAATCTGCC TGCGCAATCA AGGCATATGA GTCTAAGAAT AAGGCATCTC CAGCCAAAAT	7560

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GGCCATAGCT TCACCGAATT TCTTGTGATT GGTTAACCGC CCTCTTCGAT AATCGTCATC	7620
ATCCATAGCA GGAAGGTCAT CGTGAATCAA GCTCCCTGTA TGAATCATCT CTAAGGCAGT	7680
AGCTACCTGC GCGTGAGCAG GTTGATGGT AACCTGCAAG GCTTCCAGAA CTTCTAACAA	7740
GAGAAAAGGC CGAATACGCT TGCCACCAGC ATGAATAGAA TAGAGAACAG ACTCCC GTAA	7800
ACTAGAGGCA AACTGCTGGT CTCCATAAAA ATCTTCCAAA GCCGACTCGA CAAGAGCTAA	7860
TTTTTCTTGC TTTTCATTC AAAATCACTT TCTGTTCCGT CTTCTTGAT GACCTTGACC	7920
AAGGTCTTT CAGCCTTGTGTC CAGCGTAGCT TGGAGCTCTT TTGACAAGAC CATGCCCTTT	7980
TGAAAGGCAG TAATCGCATC TTCCAGAGCA ATTTCACCAT TTTCCAAACT TTGGACAATG	8040
GTTCAGTT CTGCTAGATT TTCCCTCAAAT TTCTTTGTT TTGACATCTT TAACCTCTAA	8100
TTCTACTTGA CCATCTCGCA TCAAAAGCGT TACTTGGTCT TTTTCTTCA AACTCTCAAC	8160
CGAATCTACA ACGGACTCTT CTTTTTGAC AATAGCATAA CCACCGCGCA CGATT CGGCT	8220
AGTATCCAAC ATGAGCAAAG CTTCCGAAAG TCGCTTGCC TCAGCAACCT TGGCGTCATA	8280
AACTAACGCC ATTGGTCTAC CTAAGAGCTT GTCCAACCTGT CCTAAACGGT CTTGATAGCG	8340
TTGGATTTG GTAACAGGTG ATAATTGTAC TAATTGATGA GTTCTTGCTT GAACTAATTG	8400
TTTGTATCA GAAATCCGAG TTCCGAAACT TTGTTTCAA CGCAGTTGCA GTTGGTCAA	8460
GCCTTCCAAA TAACCGTCAT ACAAGCGCTC AGGTTGCTA AAGATAACAG ACTGACTGCA	8520
TTTTTCTTCAA GCCTCTTGTGTT TCTTAGATAG AACATTTCCG ACTGCCGTTA CCATCCGTT	8580
TTCCCTGATTT TGCAAATGAG CTAATACATC CAACTTGGTC ACAGGTGTTG CCAGTTCA	8640
CGCCGCTGTT GGCGTTGAG CGCGTCGATC TGCCACAAAA TCTGCCAAGG TCACATCCG	8700
CTCATGCCCAAC ACTAGAGA TAACTGGCAA ACGAGATTCA AAAATAGCTC GTACCACAAT	8760
TTCTTGTGTTA AAGGCCAGA GATCCTCAAT AGAACCCACT CCACGACCAA TAATGAGCAA	8820
ATCCAATCG TCCCGTTGAT TAGCACGCGC AATATTTCTA GCAATTTCTT CGCGAGCCCC	8880
TTCACCTTGA ACCTTGGTCG GATAAAGAAG GATGTCAACA CCTGGGAATC GCCTGCTGAC	8940
GGTCGTGATA ATATCTGAA TAACGGCTCC ACTACGGCTG GTTACTACAC CAATTCTCTT	9000
AGAAAATTGG GGCGAGAGCTT GCTTGAAGCG TTCTTGAAAC AGGCCTCTT CTGTCAATT	9060
TTTCTTAAGT TGTTCAAACG GAATCGCAAG CGCCCCAACCC CCATCAGGCT CAGCTTTTC	9120
AATGATGATG GAGTAGCTAC CACTTGGTTC ATAGACCTGT ACACGCCAA TCACATTGAT	9180
CTTCATTCCT TCTTCCAGGT CAAACCTAA TTTCTGATAA ATCCCAGACCG AGATGGTCGC	9240
TTGAATAACT GCATGGTCAT CCTTTAGGGA GAAATATTGG TGAGTAGGTC GTTTACGAA	9300

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GTTGGAAACT TGACCAAGTTA AATAGACCCG TTCCAAGTAT GGGCTTTAT CGAATTCAT	9360
TTTCAGATACT TTGGTCAAAG TTGTTACCGA TAAATACCTT TCCATCTCCA CCTACTATTG	9420
ATTTACTTGC TCTTCATGG GTATTATTAT ACCAAAAATA TGCCCTAAAAA TCTCCATTAA	9480
TGTACCATTAA TGAGGGAAAA ATAGAAAAG GAGGCAAGGC CTCCACATGT GATTATTG	9540
TGTTTCGAGC TTCTTCCAAA ATCTTGCAA TCTTGGTCGT CAACAGGTCG ATAGCCACGG	9600
TATTGCTAAC CCCTTCAGGA ATGACGATAT CAGCATAACG CTTAGTTGAC TCGATAAACT	9660
GGTGGTACAT TGGTTGACC ACACCTAAGT ACTGGTTAAT AACGCTATCA AGGCTACGGC	9720
CACGCTCCCT CATATCACGC TTGATACGAC GAATAATGCG CACATCGTCA TCCGTATCCA	9780
CAAAATCTT GATATCCATC AAATCGCGCA GACGCTTGTC CTCCAAGACC AAAATACCC	9840
CAACGATAAA GACATCTTGA GGTTCTGAC GATAGGTCTT GCTACTCCGT GTATGCTCTG	9900
TATAGTCGTA GGTGGGGATG TCCACCGGAC GCCCTGCCAA CAATTCCCTTA ATCTGCTCGA	9960
TCATCAAGTC TGTATCAAAG GCAAAAGGAT GGTCAAGTTT GGTTTGACG G	10011

(2) INFORMATION FOR SEQ ID NO: 77:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 5365 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 77:

CGTGTGGTCT TAAAAATAGA AGACAAAGAA CAAACTGTG GAGGCTTTGT CCTTGCAGGC	60
TCAGCCCAAG AAAAACCAA AACAGCTCAA GTTGTGGCTA CTGGACAAGG TGTTCGTACC	120
TTGAACGGTG ACTTGGTTGC TCCAAGTGT AAAACTGGAG ATCGTGTCTT AGTTGAAGCC	180
CACGCAGGTC TTGATGTCAA AGATGGCGAT GAAAAGTACA TCATCGTAGG CGAcTAACAT	240
TTTGGCAATC ATTGAGGAAT AGAAGGAGAA AGTAAGTATG TCAAAAGAAA TTAAATTTTC	300
ATCAGATGCC CGTTCAGCCA TGGTTCGTGG TGTCGATATC CTTGCAGACA CTGTTAAAGT	360
AACCTTGGGA CCAAAAGGTC GCAATGTCGT TCTTGAAAG TCATTCGGTT CACCCCTTGAT	420
TACCAATGAC GGTGTGACCA TTGCCAAAGA AATCGAATTG GAAGACCATT TTGAAAATAT	480
GGGTGCTAAG TTAGTATCAG AAGTAGCTTC TAAAACCAAT GATATCGCAG GTGACGGAAC	540
TACGACTGCA ACAGTCTTGA CCCAAGCTAT CGTCCGTGAA GGAATCAAAA ACGTCACAGC	600
AGGTGCAAAT CCAATCGGTA TTCGTCGTGG GATTGAAACA GCAGTTGCCG CAGCAGTTGA	660
AGCTTGAAA AACAAACGCCA TCCCTGTTGC CAATAAGAA GCTATCGCTC AAGTTGCAGC	720

629

CGTATCTTC CGTCTGAAA AAGTGGTGA GTACATCTCT GAAGCAATGG AAAAAGTTGG 780
CAAAGACGGT GTCATCACCA TCGAAGAGTC ACGTGGTATG GAAACAGAGC TTGAAGTCGT 840
AGAAGGAATG CAGTTTGACC GTGGTTACCT TTCACAGTAC ATGGTGACAG ATAGCGAAAA 900
AATGGTGGCT GACCTTGAAA ATCCGTACAT TTTGATTACA GACAAGAAAA TTTCCAATAT 960
CCAAGAAATC TTGCCACTTT TGAAAGCCT TCTCCAAAGC AATCGTCCAC TCTTGATTAT 1020
TGGGGATGAT GTGGATGGCG AGGCTCTTCC AACTCTTGTGTT TTGAACAAAGA TTCTGTTAAC 1080
CTTCAACGTA TAGCAGTCA AGGCACCTGG TTMTGGTGC CGTCGCAAAG CCATGCTTGA 1140
AGATATCGCC ATCTTAACAG GCGGAACAGT TATCACAGAA GACCTTGGTC TTGAGTTGAA 1200
AGATGCGACA ATTGAAGCTC TTGGTCAAGC AGCGAGAGTG ACCGTGGACA AAGATAGCAC 1260
GGTTATTGTA GAAGGTGCAG GAAATCCTGA AGCGATTCT CACCGTGTG CGGTTATCAA 1320
GTCTCAAATC GAAACTACAA CTTCTGAATT TGACCGTGA AAATTGCAAG AACGCTTGGC 1380
CAAATTGTC GGTGGTGTAG CGGTTATTAA GGTTGGAGCC GCAACTGAAA CTGAGTTGAA 1440
AGAAATGAAA CTCCGCATTG AAGATGCCCT CAACGCTACT CGTGCGAGCTG TTGAAGAAGG 1500
TATTGTTGCA GGTGGTGGAA CAGCTCTTGC CAATGTGATT CCAGCTGTG CTACCTTGG 1560
ATTGACAGGA GATGAAGCAA CAGGACGTAA TATTGTTCTC CGTGCTTGG AAGAACCCGT 1620
TCGTCAAATT GCTCACAAATG CAGGATTGAG AGGATCTATC GTTATCGATC GTTTGAAAAA 1680
TGCTGAGCTT GGTATAGGAT TTAACGCAGC AACTGGCGAG TGGGTTAAC A TGATTGATCA 1740
AGGTATCATT GATCCAGTTA AAGTGAGTCG TTCAGCCCTA CAAAATGCAG CATCTGTAGC 1800
CAGCTTGATT TTGACAACAG AAGCAGTCGT AGCCAATAAA CCAGAACCGAG TAGCCCCAGC 1860
TCCAGCAATG GATCCAAGCA TGATGGCGG GATGATGTAA GCTTTCTATA GAAAACAAC 1920
TATAAAAAC ACAAAAGGAG GGAATGACTA ACCCTCTTT TTATAGGCTC TTTGTCAACT 1980
GTAGTGGTT GAAGTCAGCT AAGCTCGAGA AAGGACAAT TTCTGCTTT CTTTTTGAT 2040
GTTCAAAGCG ATAAAAATCC GTTTTTGAA GTTTCAAAG TTTCGAAAAC CAAAGGCATT 2100
GCGCTTGATA AGTTGATGA GATTATTGGT CGCTTCCGGT TTGGCGTTAG AATAGTGTAG 2160
TTGAAGGGCG TTGATAATCT TTTCTTATC TTTGAGGAAG GTTTAAAGA CAGTCTGAAA 2220
AATAGGATGA ACTTGCTTAA GATTGTCCTC AATAAGTCCG AAAAATTCT CCGGTTCTT 2280
ATTCTGAAAG TGAAACAGCA AGAGTTGATA GAGCTGATAG TGATGTTCA AGTCTTGTGA 2340
ATAGCTCAAAG AGCTTGTCTA AAATCTCTTT ATTGGTTAAA TGCATACGAA AAGTAGGGACG 2400
ATAAAATCGC TTATCACTCA GTTTACGGCT ATCCTGTTGT ATGAGCTTCC AGTAGCGCTT 2460

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GATAGCCTTG TATTCATGGG ATTTTCGATC CAATTGGTTC ATAATTTGAA CACGCACACG	2520
ACTCATAGCA CGGCTAAGAT GTTGATACAAT GTGAAAGCGA TCCAACACGA TTTTAGCATT	2580
CGGGAGTGAA ACAGTCTGGG AGACTGTTTC AGCCTGAGCC TAGAAATTG AAAGCGAAC	2640
TGTTTAGCCA AGTCATAGTA AGGACTAAC ATATCCATCG TAATGATTT CACTTGACAA	2700
CGAACGGCTC TATCGTAGCG AAGAAAAGTGA TTTCGGATGA CAGCTTGTGT TCTGCCCTCA	2760
AGAACAGTGA TAATATTAAG ATTATCAAAA TCTTGCACAA TGAAACTCAT CTTTCCCTTA	2820
GTGAAGGCAT ACTCATCCCA AGACATAATC TTTGGAAGCC GAGAAAAATC ATGCTCAAAG	2880
TGAAAGTCAT TGAGCTTGCG AATGACAGTT GAAGTTGAAA TGGCCAGCTG ATGGGCAATA	2940
TCAGTCATAG AAATTTTTTC AATTAACCTT TGAGCAATCT TTTGGTTGAT GATACGAGGG	3000
ATTTGGTGAT TTTCTTTAC CAGGGGAGTC TCAGCAACCA TCATTTTGA ACAGTGATAG	3060
CACTTGAAAC GACCCCTTCT AAGGAGAATT CTAGAAGGCA TACCAAGTCGT TTCAAGATAA	3120
GGAATTTTAG AAGGTTTTTG AAAGTCATAT TTCTTCAATT GGTTTCCGCA CTCAGGGCAA	3180
GATGGGGCGT CGTAGTCCAG TTTGGCGATG ATTTCCCTGT GTGTATCCTT ATTGATGATG	3240
TCTAAAATCT GGATATTAGG GTCTTAATA TCGAGCAGTT TTGTGATAAA ATGTAATTGT	3300
TCCATATGAA TCTTCTAAT GAGTTGTTT GTCGCTTTTC ATTATAGGTC ATATGGACT	3360
TTTTTCTAC AACAAAATAG GCTCCATAAT ATCTATAAGG GATTTACCCA CTACAAATAT	3420
TATAGAGCCG AAAATTCAAA TCTAATATAT GCAGACTACT TTGAAATGAA ATTAAAAAAA	3480
TTATTTAAAGG ATGACACAAA AGTTTTGAA AAATCTACAT TCAAATTGTT AGAAGGATAT	3540
AAAATATACC TGACAGAAC TAAAGAATCT GGAATTAAAC AAATGGACAA TGTCATAAAA	3600
TATTTTGAGT TTATTGAATC TAAAAGTATT GCTTTATATT TTCAAAAACG ATTAAATGAG	3660
CTGATAGATT AAATAGCATT TTCTCTGTT AGATATTGTT TTTAAATAT TGTACTAAAT	3720
GATTGATGCT ATGTGAAAT ACAAAAAAAT GTTTTGATA CGAAGTGAC CTGTATTTT	3780
TATACTAATC ATTTTCGTAT TTTTGTTATT AAACGATATA AGTTTGTTGT AAACCTACAA	3840
GGAATAAAAGA CATTAAAAAA TAACAGTATA TCTATTTGTT TTATATATTT TACGAATTCT	3900
GCATAAAATCT CTTCTAGTA ATGTGTTGTA ACTCTGCTAT AATAGATTTA TTCTCTTTG	3960
TGTTTACACA ATTTATTTTA TAGTACCAA AAAGGTCAGG ATTTGTTCC TGACCTTTGA	4020
CAACTTACCC GATTCTTAG TTCTACATAG CGCTTGTACC AAATGTTAC ATAGGCTTCT	4080
GAGAAAGGAC CACGTC CATT GTTAATCCAA TCAACAAGAA TTTTGACATG TTCTTTAAA	4140
ATATAGTCCA AGTCATCAGA ATAATTCATT TTGCGTTGT GACGCTCGTA CTCTTCAACG	4200
TCCAAGAGAC GTTTTCCCC ATCTGAAAA ATTTAACAT CCAAATCGTA ATCAATATAC	4260

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TTCAGTGCTT	CTTCATCCAG	ATAGTAGGGG	CTAGCCATAT	TGCAATAGTA	AGAAGTTCCA	4320
TTATCACGAA	TCATGGCAAT	GATATTAAAC	CAATATTCT	TGTGAAAGTA	AACAATAGCC	4380
GGTCTCGAG	TGACCCAACG	ACGACCACATCA	CTTTCGGTAA	CAAGTGTATG	ATCGTTGACA	4440
CCAATAATGG	CGTTTCTGT	TGTTTTAGT	ACCATGGTGT	CCCGCCAAGT	TCGGTGGAGA	4500
CTCCCACAT	GCTTATAACT	TTGAATTGTA	ATAAAGTCGC	CTTCTTTGG	AAGCTTCATA	4560
ACTAACCAAC	TTTCTACAAT	TTATAAGTTT	ATCATTACT	ATTGTACCAT	AAAATTACCC	4620
AAAATCTGT	AATTCACTT	GGAAATATTA	AAGATATTCT	CTAAGAGCGC	TTGCTATATC	4680
CGAAAAATCG	TAGCCCTTC	GTGCTAAAC	TTGAGTTAA	CGCTGCTTC	GTTCTATCC	4740
TTCATACTT	CGGGCATACT	TAGTATATTG	CTTATCAACT	TCCTTGAAGA	TGAGTTCC	4800
AGTCGTTCT	TCATCAACTT	GACTATCAA	TTCGTCAAAG	GCAATTAG	CATCAAAATA	4860
AGAGAAGCCC	TTGTTAGTCA	AGTTCTGGAT	AATCTTATCT	TGCAGGGCAC	GAGCTGGAAG	4920
TTTCCCTCA	TATTTTTCTA	ATAGTTTATT	GGCTACACGT	TGAGCAACTT	CCGAAAAATC	4980
AAAATCATTC	AAGATTTCTT	CTATAGTAGA	TTTTGAAATT	CCTTTTGTC	CTAATTTCTG	5040
AGTCAGTACA	TAAGGTCCCT	TGTCTCCTGA	AAGTTGATTG	GCATTGATGA	TAGCATAAGC	5100
GTACTGGCTA	TCATTAATCC	ACTTCTCTTC	TTTAAGATTA	GCAATGACTT	GAGAAACGAT	5160
GTTTTCATTA	ATATCATATT	TTTCAGATA	TTCTCTGACC	TCTTTTTCAG	TACGTGCTT	5220
AAACGTTAAC	TGGTAGAGGG	CCAGATTCTT	ACCATAAGAA	AATTGAGCAA	AGTCTTGAAT	5280
CTCTTCAAT	TCCTCTTCGC	TTATCACCTT	ATCTCTCGAT	AACATAAAAC	GAACAATTGT	5340
GTCTCGGTG	ATATAGCATT	TGTCG				5365

(2) INFORMATION FOR SEQ ID NO: 78:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 3636 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 78:

TTTCCAGAAA	GAAGTTGAGT	AAAGTCTTAA	TCAAAGAGAA	TGACTTCCGT	ATTGGAAC	60
ACATTAGTT	TTATTTCTAC	TTTACTAGCG	TCCGCCCTAG	CATTTCTAA	ATCTTTAAC	120
TCTTCTGTTG	CCCTATTTAT	AGCCAGCTGA	ATAACTGCTT	GAGGATTTTC	ACTCAGTCCA	180
TGAAGCTTAT	CGTCCACCGA	AGTATAAAGA	CTCGAATGCA	TGACTTGTAA	AATAATCAGA	240

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GTCATTGTAG	AAAAAATCG	GGTGAAGACA	CCGAAGTTGC	GGATAAAATA	ACTAAAGTCA	300
TCCGCATACC	ATGTTTTTT	AAGTTTACTG	AACATCTTT	AAAAGATACC	CAACACTACG	360
CAAAGTTGC	AAATTCTCTG	CAAAGTGGT	TCCCTTAAT	TTCTTACGGA	CTTTGAAAC	420
ATAGACTTCG	ACAACCGAA	TCGTTGTATC	ACTATCAAAT	CCCCATAGAC	GGTCAAAAAT	480
CTGCGTCTTA	GGCAAAATCA	CATTTGATT	TTGAAGGAAA	TAAACTAGTA	AATCGAACTC	540
TTTCCCCAGC	AATTGACAG	GAGTATCTTC	AACTTTAACG	GTATTGGTTG	ATAAATTAAC	600
CACGATATTC	CCATAAGTCA	AGGTGTTTC	ATTAACATTC	CCTGAACGTT	TGAGAAGGGC	660
CTGAATCCGC	ATTTTAAGTT	CTTCTAGGTA	GAAAGGTTTG	GTCAGATAAT	CATCCGCTCC	720
CAGTTCAAAT	CCATGTCCT	TGTCATCCAA	ACTTTCTTG	GCAGTCATAA	TCAGAACTGG	780
TGTCGTAATT	CCCTTTAC	GCAATTCTTT	TAAGACTTG	AAACCATTAA	TTTCTGGCAA	840
CATCAAATCC	AGCRAAAATCA	AGTCATAGAC	ACCACTCTCA	GCTTCTAGA	GACCTTCTTC	900
TCCATCAAAT	ACCTGCATAA	CATCCGAAA	ATCGTCTAA	AAAGTCAAAATA	CTGAATTGAA	960
CAGACCTAGG	TCATCCTCAA	CCAATAAGAT	TTTATCATG	AGAAAATCCT	CCTTATTAAA	1020
ACTATTATAC	CAAATTGCC	TTAAAAAAA	CTCAACTCTC	TGCATTTAC	ATGAGATAGC	1080
TGAGTTTCT	TTTATTTTA	GGCTTATTAA	TGCATTTCCG	TATTGAAGAA	CAACTGCTTC	1140
GAETGCAGCT	TTTCACGGC	TAATCAAGTC	AACACGCGCT	GCAATTCCCT	TGATTCCCAT	1200
ACCGATGTTA	CGGCTAAGAG	CAAGGTAGA	AAAGTCAGA	TCAAAGAACT	CCTTGTATTTC	1260
CGCCAAGCGT	TGCTGAGTCT	AAATACATG	AGCAGGAAGG	ATAACAAAGC	TATCAAAGCT	1320
CATATCTCCT	CCAAGGGCTG	CCTTAATCCA	AGCCCAGTTT	TCACCGGCC	AAGACCAAGC	1380
TGTTTCTGA	GTTGCTTGAT	GAGCTAGGAA	TTGGTAATAC	CAAGCAGACA	AGTCCTGTGG	1440
TTTGACCACA	AATTGTCCT	TCCAAGAAGT	AATCAGGTTT	TGGATATTAT	CCGCATCTGT	1500
ACTGTATGCA	AGAGCTGCTG	CCAACTGGCG	TTAAAGACA	GCATCTGTTG	CGTGACTATA	1560
AGTATCAAGA	TAAAGTGCTA	ACAAGTCTTT	AGTCTCATGA	TGTTTCATCT	CATTAATCAG	1620
AACTTGTGAG	CQAATAGCTG	CTGGGAGTCC	TGCAAGATTC	TCCTTGTGTG	TTGCGAAGAT	1680
TTGGCTAGCG	ACTTGACTAG	CTTCTGCATC	ATTGAGCGA	ATCATCATCG	AAACAGCCAG	1740
CTGACGAACC	AATTCTCCT	CATCTGATTC	TCCGTCTTTA	GCTTCAAAAC	CAAGACGGTC	1800
ATAGTTATGA	CGAGCCAATT	TAGCAACCAG	TCCTTGAAG	GCTGTTTCAG	CATCCGTTCC	1860
TTCATCAATA	AAGCGCTCAA	GGGCTGAAAT	CACTTGAGAA	ACAGCTGAAA	CCACCAAGATA	1920
AGACTCTTCC	TTAGCAAGTT	TATCAAGAAC	TGGAAGCAAG	TCTGCATAAG	AAATGTGCC	1980
TGCCTCAGCC	AACAAACGAC	GTTCTTGAAC	AATTGCAAGT	TTGCTTGTGT	TATCAAGTGT	2040

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CTCTAGCTCA GCAAGAACAG CTGCTAACAA GTCTCCTTGA TAGTCGGTAA TATAGTGGC	2100
AGTATTTCA GTGTTGAGAC GAAGAGCTCC TTCATTTCA GCAAGAAGAG CTGCGTAGCC	2160
AGGGATTCG ATACTTCAG TTTCGAGTGT ATCAGGCAAG CCTTCCAGT TGCTATTGAG	2220
GGGCACCACC CAGAGACGGT TCTTGCTTC GTTCTCACCG ATGAAGAATT GTTTTGTGA	2280
AATCTCAAG ACATCATTTC CAACTTTAAC AGTAAGAACT GGGTAACCAG GCTGTTCAA	2340
CCAAGAATCC ATGAAGGCTG CGACATCACG TCCTGACGCT TGACCAAGGG CATCCCAAAG	2400
GTCACTACCA ATGGTGTGCG TGTATTGGTG TTTTCAAG TAGGCGTGC AACCTTTAGC	2460
AAAATCAGCA TCTCTAGCC AACGGCGAAG CATGTGCATG AGACGGCTTC CTTTGGCATA	2520
GACGATAGCG CCGTCAAAGA GTGTATTGAT TTCATCTGGA TGTTTAACCT CGACGTGGAC	2580
AGACTGAACG CCATCAGTAG CGTCACGTT AAGAGCAAGA GGTACTCCAC CTGTTGGAA	2640
ATCTTCAAAG ATATCCAGC TTGGTTCGAT GGATTCACA CAGACGTATT CCATCATATT	2700
AGCGAAACTT TCATTGAGCC AAAGGTCATC CCACCATTTA ATAGTCACGA GGTTCCAAA	2760
CCATTGGTGA GCCAATTCA GGGCCACAAC AAGGGCAACT TGTTGACGGC TAGCAAATGT	2820
AGAGTTCTCA TCGACAACCA AGTAAACTTC ACGGTAGGTC ACAAGACCCC AGTTTCCAT	2880
AGCACCAGCT GAGAAGTCAG GAAGGGCGAT GTGGAGAGAT TGAGGAATTG GGTACTTAA	2940
TCCATAGTAA TCTTGTAAA ACTCGATAGA GCGAACAGCG ATATCCAGTG AGAAATCAAG	3000
ATTTGAAGT GGATGTGCTT TGGTTGAGTA GACACCTACC AGGGTACCAT TTTTAGTTT	3060
AGCGGTCACC CCTTGCAAAT CACCAGCAAC AAAGGCCAAC AAGTAAGAAG ACATGCGAGG	3120
TGTTGCTCTCA AACTTCCAGA TACCTGTTTC CTACGGTTT TCAACATCGA TTTCTGGCAT	3180
GTTTGACAAG GCCAATTCA CTTCTGCTTG GTCAAAGCGA AGAGAGAGGT CAAAAGTTGC	3240
TTTGGCTTCA GGCTCATCCA CACATGGAA AGCTTCGCGC GCAAAATGGC TCTCGAACTG	3300
AGTAGACAAG ACCTCCTTCT TGACTCCATC AACTGTATAA TAAGAAGGGT AAATCCCTGT	3360
CATGTTGTCT GTAATTTAC CAGAAAAGGC AAAACCAAT TCAACTTGAC CAGCCTCAGC	3420
CAATTGATA TGAAGGGCTT CATTGTCATG GTCAAACGTGA AATGGACGAG CTTGACCTGC	3480
AACTTCTACA GAGGTGATTT CCAAATCTTT TTGGTGGAGG GAGATGCGGT CACTCTGTGC	3540
TTGACCAGTG ATGGTCACTT TCCCAGAAAA AGTCTTGGTC TCACGACTCA AATCTAAAAA	3600
TAAATCATAA TGTCAGGAA CAAATTGCTT AATGGG	3636

(2) INFORMATION FOR SEQ ID NO: 79:

- (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 5066 base pairs

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- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 79:

ATAGCGTGTA ATAATCGATT TTAGAGGTAC CATAAGCCAC CTCCTACAAA TAGAAACCGA	60
TATAAAATCAA TGCCTTCCAC CCTTAGACTT CCCTAGTTCC TGTCTCAAGC GAAACATTTC	120
TTTGAAACAG GAATAAGTTA ACCAATTCA ACCAATAGCT AGCAGAATAA AAAGAAACCA	180
AATGCCCAT AACTTGATAT CTGTCACATT TCTCAAGACG GTATTGAAAA ACAGAACTGA	240
AAACAATGTC CAAGCAAGGC TAAAAAGAGA ATAGAAGGGG ATGTAAAACC AGTAAAATA	300
ATAAAAAAATT GGAAAAAAACT TACTATTCT GTTGGCCTTT TCAATCCAGT TATCAAATA	360
AAAGTACGGT GCTAAAAGTA AGAATTAAA CAAATGTTCC ATCACCGACA TCCCCCTTC	420
TTTTGATAGC GTTTCTATT ATTTTATTAT ATCAAAAAAA TCCGGAACGT TCATTCCAGA	480
TTCTACTTTT TTATTGCGT TTTCTTGCAG TGAGATGAAT CGGTGTTCCC TCAAAACAA	540
AGGCCTTGGCG GATTGATTT TCCAAGAAC GCAGGTAAGA AAAGTGCATG AGTTCTTCTT	600
CATTGACAAA GATGACAAAG GTTGGTGGTT TGTTGCCAC TTGGGTCGCA TAGAAAATCT	660
TGAGACGTTT TCCTTGTCT GTGGTGTG TGTTGATGGC AATGGCATCC ATGATGACAT	720
CGTTCAAGAC AGCTGATGGA ATACGTGTAT TTTGACTTTC GCTGATTGC TTAATCATCT	780
CAGGAAGTTT GTGGAGACGT TGCTTGGTTA AAGCTGATAC AAAGATAATC GGTGCGTAAG	840
GCAGGTATTG GAACTGCTCA CGGATATCTT CTCCTCAGTT TTTCATAGTG TGTTATCTT	900
TTTCAAGCGT ATCCCACITG TTGACCACGA TAATCATCCC TTTACCAAGCT TCATGGCAA	960
ATCCTGCGAT ACGCTTGTG TACTCACGAA TGCCTTCTTC CGCATTGATG ACCATCAAGA	1020
CCACATCTGA ACGGTCATAA GCACGCATGG CACGCATAAC AGAGTATTTC TCAGTATTTC	1080
CATAAAACCTT ACCAGACTTA CGCATACCAAG CCGTATCAAT CATGGTAAAC TCTTGACCAC	1140
CTGTATCTGT AAAGTGGGTA TCAATGGCAT CACGAGTTGT TCCAGCAACA GGACTAGCAA	1200
TAACACGGTC TTCTCCAAG ATAGCATTGA TCAAGCTTGA TTTTCCAACG TTAGGACGAC	1260
CAATCAAGCT AAACTTAATG ACATCTGGAT TTCTCCCTC ATATTCATT GGAAGATTTT	1320
CTACGATCGC ATCTAGCACA TCCCTGTAC CGATTCCATG GACAGATGAG ATAGGCAATG	1380
GTTCACCCAA ACCGAGAGCA TAGAAATCAT ATATATCATT TCTCATCTCA GGGTTGTCCA	1440
CCTTGTGAC TGCGAGGATA ACTGGTTTGT GGGTCTTATA AAGCTTACGA GCTACGTATT	1500
CGTCTGCATC AGTAATTCT TCCTTACAG ACACGACAAA AACGATAACA TCTGCTTCTT	1560

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CCATGGCAAT TTCTGCCTGG TGCTTGATTT GTTCCATGAA AGGAGCATCG ACATCATCAA	1620
TTCCTCCTGT ATCAATCATG CTAAAAGAAC GATTGAGCCA CTCACCCGTT GCATAAATAC	1680
GGTCACGTGT CACTCCTTCG ACATCTCTA CAATGGAGAT TCGCTCACCA GCGATCCGAT	1740
TAAATAGGGT TGATTTCCC ACATTGGGAC GTCTACAAT GGCAATAGTT GGTAGGGCCA	1800
TAATTTCTCA CTTCTACAA TAATTTCTTC TGTTCAAGAT TTTTTCTAGT TGAGCTTGTT	1860
TCAGCTTGAC CAAACTGTTG TGCTAGGCAGC TGACTCCAGC TTGTGGTCGC ACGCGCCCCA	1920
GCATAGTCAG CCTGAACACG GTCATAAGCT TGGATTGCCT CAGTTGACTG TTCTTGGTAT	1980
TCTTCCTCAA AGACAACATT CTCTAGTGGC AGTCTCGGTT TCATATCATG ATGTTGATTT	2040
GGCACACCCA GTGCCATCCC AAAGACAGAA TAGGTGTAGT CAGGTAGGTT AAAGAGCTCT	2100
GCCACTTCTT CAGACTTGTA TCGAACCAAA CCGATAATCA CACCACCCATA GCCCAAGCTT	2160
TCAGCTGCCA ACAAGGCAGT TTGTCCAGCA AGAGCTGCAT CGACCGAAGT AATCAAGAGA	2220
CCTTCCACAC CTTGGGTTG GAAGGTGTG GTATGAAGTC GGGCTCCCTT TTCTGCTCGG	2280
TTCAAATCTC CGACAAAGAG AAGGAAAACA CGAGACTGGC GAATGGCTTC TTGAGGTACC	2340
AATTCAACCA AGGCATCTTT CTTCTCTTGA CTTCGTACCA CAATCACAGA GTAGGATTGG	2400
AAATTCTTCC AAGATGATGC CATCTGGCT GCTGTAAAAA TCTCATTTAA GTCTACTTGG	2460
GGAATTCTT GCTCTTTAAA CCTGCGCACT GAAGTATGAG CCTTCATCAA TTAAATGGTT	2520
TCTGTATCG ACGGTTTACT CCTTCTAAAC GAGTCTCCTC AGCCAAATAA CGGATGCGTT	2580
CCATGACCCG TCTGGCTTCC CAGGTTTCGT CATTTCATG TTTCACTTTC GCAAAATGCT	2640
TCTCCAAATC TTCAAAGTTG AAGTTGGATG TGAAAAGGT CGGTAATTT TCCTGCATCC	2700
GATATTGGAG AATGACCTGC AGGATTTCGT CACGCACCA AACGGTTGAT TGCTCGCGC	2760
CAATATCATC TAAAATCAGG ACCTCAGACA CCTTAATCTC ATCCACCAAG GTCTTAACAT	2820
TGCCATCACT GATAGCATTT TTGACATCAA TGACAAAGCT AGGATAGTGG AGGAGAGTTG	2880
ATGAAACACC ACGTTTTCT GATAAAATCAT GAGCTAAGGC CGCCACCATG AAACTTTAC	2940
CCACACCAAA GTCTCCATAT AAGTAAAGAC CTTTTCGAAT AGCTGGATAT TGCTCCACGA	3000
AGGCTAGTAG CTTTCTAAA ACTGGTAAGC GCCCCAAATC ATCCAAGTCA ACTTGAGCCA	3060
AACTAGCTTT CTTGAGACTG GCTGGTAGAT TGATTAACCT GAGACGGTTC TPAATAGCCG	3120
CTTCTTTTC AGCCCGCATT AGCTCAGGAG TTTCTTCATA TGAAACATCT GCATAACCAT	3180
GATTCTTAAC CAAAATCGGC TTGTAGCCTT TGGCAATATA ATCCGTATCC CCACGGAGAA	3240
ACTTGTCACG CTCGGTGATG TACTGATTAA ACTTGGAGAT ACTGCGATTT AATTCTTTG	3300

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GAGTTAAGGA TTCTTGCTGG ATAAAGGCCG CAACATCAGG GTCCTTCATG ATTTTCTGGA	3360
CCAAATCTG ATAATAAAAA CGGCTGGTT GACCTTGAG TACGTCTCCG ACACTTTCCA	3420
TCTAACCTCC TCCTTTTCT AATCGAGCTA ATACTTCTTG CTTCTTACGT TCTAGTTCCA	3480
GACGAGTTTC CTCGCTGGTT TCATTCTTAT ATTCAAGGATT ACTCCATTAA GGAACATTGG	3540
TTTTTCTGGG GGCAGTCTGA TTCTGTTTT GTGTTTTGC TTTCTGCCCT CGATCACGAA	3600
TCGTAACAC GGCTCTTCT GCCAATGAA TCTTTGATA GGCATAGTCA TTGGCTACCT	3660
TCATGGCATA TTTCTCATG ATATTGCGC AATCCACCTT ATTAAGGTC AATAAGAGAA	3720
TAATATTGAT GACTTCGTCC AGTAAGCCCA AGCCAGCCAT CTGTTGCAAG AGTTCTCTT	3780
CTGTTGGGT AATGGTCCC TTGCGTGTGTT GCTTGATTTC TGCTAAGAAC TGCAGGGCAG	3840
TTTTACTTTT AGCTTCTTG ATAATGGTCG CTTCCTTAAG ACTAAAGTCA GAGGAAACTG	3900
GTTTTGAGC AATTTTTCA CGCATGCGTT TGTTGAAAT AACCTGGAA ACAGCTGTTG	3960
ACTTGGCCAA TTGATAGGTT TCAAACCAAG TCCATTCTT CTCCTCGGCA ATAGCAAAGA	4020
GGTTTAAGAC ATCGGACTGC TCATCCGCAA AACGAAGTCC ATCTCGAGCC ATCAGCTGGC	4080
GAAAATGTTCA AAGTCAAAA TCATTGGCCA CTTCTCTT GAGACCAAGG TCTTCTTGAC	4140
TGCCTAGTTC TGCCAATTCT GGAAAGACTT GATTGAGTGA GACAGGTATT TCTTCACCCT	4200
CAGCACTTTC AACTTCAAA TCCTCCACAG CTACATGCC AATCTTTTTC TCTAAGAGTC	4260
TCCGATAAAC AGGATGCCA AAGAAGTCTT GACTAGATAG AGGAGCATGG AGGGCTAGCT	4320
GATAAACATC ACCCTTTGA TAGAGGGTCA AGAGATTTAA AGCAGATAAG ATTTTCAATG	4380
ATTTTATCG TCTATCCATC CCAAAGTTGA GATGGTTGAG AATGCTTGAA AAAAGATATT	4440
CCTTCTACC ATTATCCAA AAACGTGATTG TATAAAGATA AAGGCTCAGT GCCTCCTGAC	4500
CGATAATCGG GAGGTAGCAC TGTACCAAGAG ATGAGGTATC TTGCGACACC CGATTATTCT	4560
TTAGATAAGA AAAACGGTCA ATTGGCTTCA TTTATCTTTC CTTTTCTTT TTAGAGGACT	4620
GGGTGATTTG TTGGAGCAAG CTCTCTAACT CACTGACATC CTTAAAACTA CGATAGACAC	4680
TAGCAAAACG TACATAGTA ATCTCGTCCA ATTCAGCCAA CTCCCTCCATG ACGAGTGAAC	4740
CAATGTCCTC ACTTTGAATT TCATTCTCATC TTGACCCACG GAGTTCTGT TCGATACGAT	4800
TGACTACCAT GTTGATTTCA TCACTTGACA CAGGACGTTT CTGGGCTGAG CGGATAATCC	4860
CATTAAAGAT TTTATCTCTG GAGAATTGTT CCCGTGTGCC ATCTTTTTA ACAACCACTA	4920
AGGTTCTTTC TTCTACTCGT TCGTAGGTTG TAAAACGGTG TTGGCATTG TCGCACTCAC	4980
GTCTTCTACG AATGGTGTTC CCTTCTTCTG CTTGGCAGCT ATCGATAACA CTTGACTTG	5040
TAGCCCCACA TTTGGACAG GGTACC	5066

(2) INFORMATION FOR SEQ ID NO: 80:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 9607 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: double
(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 80:

CACTTGAAGT ATTTGAAACA GCTATGGAAA ACATCATGCC TGTACTTGAA GTACGTGCAC	60
GTCGTGTTGG TGTTCTAAC TACCAAGTCC CAGTTGAAGT TCGTCCAGAA CGTCGTACAA	120
CACTTGGACT TCGTTGGTTG GTAACAATCG CTCGTCTTCG TGGTGAACAC ACAATGCAAG	180
ACCGTCTTGC AAAAGAAATC TTGGATGCTG CTAACAACAC TGGTGCAGCA GTTAAGAAC	240
GTGAAGATAAC TCACCGTATG GCTGAAGCTA ACCCCTGCATT CGCACACTTC CGTTGGTAAG	300
ATAGGATGCG AAAGCGTTAA GAAAGTCCC GAGAAAATAG GGAATCGAAG CAGGTTGCGG	360
TTGCAACCAA TGAGATTCACTTCTCC AGACTTTAG CTTGAGCTCA ACTAAATCAT	420
GATGCTAGGA ACGGTAAGGA TGCAAGGTAA AAATAGGAAA CTGACGCAGT ATTGACGAA	480
TACAAGGAGT TTTATCTTT TCACGCAGCA TCCCCTTCCA GCTCACATCG GCTAACTAAC	540
TTTAGCCCCG GTTCAAAATTA GCTAAATCGA TTAGTATTAG CTATAACTCA GCTTACCATC	600
TCGTAAGTTG AAACCAACAA TAGCATGAAA ACATTGAGAA CGGGTAGGTC CTGCCTATCC	660
GTTTTTATTA AAATCGTGT ATAATAGAAT AGAAATCAAA AATAAATAGG AGAAACAAAC	720
CTCATGGCAC GCGAATTTTC ACTTGAAAAA ACTCGTAATA TCGGTATCAT GGCTCACGTC	780
GATGCCGGTA AAACAACAAAC TACTGAGCGT ATTCTTACT ACACGGTAA AATCCACAAA	840
ATCGGTGAAA CTCACGAAGG TGCGTCACAA ATGGACTGGA TGGAGCAAGA GCAAGAACGT	900
GGTATCACGA TCACATCTGC TGCGACGACA GCTCAATGGA ACAACCACCG CGTAAACATC	960
ATCGACACAC CAGGACACGT GGACTTCACA ATCGAAGTAC AACGTTCTCT TCGTGTATTG	1020
GATGGTGCAGG TTACCGTTCT TGACTCACAA TCAGGTGTTG AGCCTCAAAC TGAAACAGTT	1080
TGGCGTCAAG CAACTGAGTA CGGAGTTCCA CGTATCGTAT TTGCCAACAA AATGGACAAA	1140
ATCGGTGCTG ACTTCCTTTA CTCTGTAAGC ACACCTCACG ATCGTCTTCA AGCAAATGCA	1200
CACCCAATCC AATTGCCAAT CGGTTCTGAA GATGACTTCC GTGGTATCAT TGACTTGATC	1260
AAGATGAAAG CTGAAATCTA TACTAACGAC CTTGGTACGG ATATCCTTGA AGAAGACATC	1320
CCAGCTGAAT ACCTTGACCA AGCTCAAGAA TACCGTAAA AATTGATTGA AGCAGTTGCT	1380

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GAAACTGACG AAGAATTGAT GATGAAATAC CTCGAAGGTG AAGAAATCAC TAACGAAGAA	1440
TTGAAAGCTG GTATCCGTAAC CGCACTATC AACGTTGAAT TCTTCCCAGT ATTGTGTGGT	1500
TCAGCCTCA AAAACAAAGG TGTCAATTG ATGCTTGATG CGGTTATCGA CTACCTTCCA	1560
AGCCCACCTG ACATCCCAGC AATCAAAGGT ATTAACCCAG ATACAGACGC TGAAGAAATT	1620
CGTCCAGCAT CTGACGAAGA GCCATTTGCA GCTCTTGCT TCAAGATCAT GACTGACCCA	1680
TTCGTAGGTC GTTTGACATT CTTCCGTGTT TACTCAGGTG TTCTTCAATC AGGTTCATAC	1740
GTATTGAATA CTTCTAAAGG TAAACGTGAA CGTATCGGAC GTATCCTTCA AATGCACGCT	1800
AACAGCCGTC AAGAAATCGA CACTGTTTAC TCAGGTGATA TCGCTGCTGC CGTTGGTTG	1860
AAAGATACTA CAACTGGTGA CTCATTGACA GATGAAAAAG CTAAAATCAT CCTTGAGTCA	1920
ATCAACGTTG CAGAACCGAT TATCCAATTG ATGCTTGAGC CAAAATCTAA AGCTGACCAA	1980
GACAAGATGG GTATGCCCT TCAAAAATTG GCTGAAGAAG ATCCAACATT CCGCCTTGAA	2040
ACAAACCTTG AAACCTGGTGA AACAGTTATC TCAGGTATGG GTGAACCTCA CCTTGACGTC	2100
CTTGTGATC GTATGCGTCG TGAGTTCAAA GTTGAAGCGA ACGTAGGTGC TCCTCAAGTA	2160
TCTTACCGTG AACATTCGG CGCTCTACT CAAGCACGTG GATTCTCAA ACGTCAGTCT	2220
GGTGGTAAAG GTCAATTCCG TGATGTATGG ATTGAATTAA CTCCAAACGA AGAAGGTAAA	2280
GGATTCCGAT TCGAAAACGC AATCGTCGGT GGTGTGGTTC CTCGTGAATT TATCCCAGCG	2340
GTTGAAAAAG GTTGGTAGA ATCTATGGCT AACGGTGTTC TTGCAGGTTA CCCAATGGTT	2400
GACGTTAAAG CTAAGTTTA TGATGGTTCA TATCACGATG TCGACTCATC TGAAAATGCC	2460
TTCAAGATTG CGGCTCACT TTCCCTTAA GAAGCTGCTA AATCAGCACAA ACCAGCTATC	2520
CTTGAACCAA TGATGCTGT ACAAATCACT GTTCCAGAAG AAAACCTTGG TGATGTTATG	2580
GGTCACGTAAC CTGCTCGTCG TGGACGTGTA GATGGTATGG AAGCACACGG TAACAGCCAA	2640
ATCGTTCGTG CTTACGTTCC ACTTGTGAA ATGTTCGGTT ACGAACAGT TCTTCGTTCT	2700
GCATCTCAAG GACGTGGTAC ATTCACTGATG GTATTTGACC ACTACGAAGA TGTACCTAAG	2760
TCAGTACAAG AAGAAATTAT TAAGAAAAT AAAGGTGAAG ACTAATCCGT CCTCACTCTA	2820
GAAGGAAGTC ACTTGTGGC TTCCCTTTGT CTTTAGAAAA TACCTCTAAA TATGGTAAAAA	2880
TAGTAGAAGA ATAATGTGAG GAAAATGAAT GTCAAATAGT TTTGAAATTT TGATGAATCA	2940
ATTGGGGATG CCTGCTGAA TGAGACAGGC TCCTGCTTTA GCACAGGCCA ATATTGAGCG	3000
AGTTGTGGTT CATAAAATTA GTAAGGTATG GGAGTTTCAT TTGCTATTTT CTAATATTTT	3060
ACCGATTGAA ATCTTTTAG AATTAAGAA AGGTTTGAGC GAAGAATTCTT CTAAGACAGG	3120
CAATAAGCT GTTTTGAAA TTAAGGCTCG GTCTCAAGAA TTTTCAAATC AGCTCTTGCA	3180

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GTCCTACTAT	AGGGAGGCTT	TCTCTGAAGG	TCCATGTGCT	AGTCAGGTT	TTAAGTCCCT	3240
TTATCAAAT	TTGCAAGTTC	GTGCTGAGGG	TAATCAGCTA	TTTATTGAAG	GATCTGAAGC	3300
GATTGATAAG	GAACATTTA	AGAAGAAC	TCTTCCTAAT	TTAGCCAAAC	AACTTGAAAA	3360
GTGTTGGTTT	CCAACATTTA	ACTGTCAAGT	CGAGAAGAAC	GATGTCCTGA	CCCAAGAGCA	3420
GGAAGAGGCC	TTTCATGCTG	AAAATGAGCA	GATTGTTCAA	GCTGCCAATG	AGGAAGCGCT	3480
CCGTGCTATG	GAACAACTGG	AGCAGATGGC	ACCTCCTCCA	GCGGAAGAGA	AACCAGCCTT	3540
TGATTTTCAA	GCGAAAAAAAG	CTGCAGCTAA	ACCCAAGCTG	GATAAGGCGG	AGATTACTCC	3600
TATGATCGAA	GTGACGACAG	AGGAAAATCG	TCTGGTATT	GAAGGGTTG	TTTTTGATGT	3660
GGAGCAAAAA	GTGACTAGAA	CAGGTCGTGT	TTTAATCAAC	TTTAAATGA	CGGACTATAC	3720
TTCAAGTTT	TCTATGCAA	AGTGGGTTAA	AAACGAGGAA	GAGGCCAGA	AGTTTGACCT	3780
CATCAAGAAC	AATTCTTGGC	TCCGAGTTCG	AGGGAAATGTG	GAGATGAATA	ACTTCACACG	3840
CGATTGACT	ATGAACGTAC	AGGATCTGCA	GGAAGTTGTT	CACTATGAGC	GGAAGGATT	3900
GATGCCAGAA	GGTGAGCGTC	GGGTTGAGTT	TCATGCTCAT	ACTAACATGT	CGACTATGGA	3960
TGCTTGCCA	GAGGTCGAAG	AGATTGTTGC	AACAGCTGCT	AAAGGGGAC	ACAAGGCGGT	4020
TGCTATCACG	GACCATGGG	ATGTCCAGTC	CTTTCCACAT	GGCTATAAGG	CGGCTAAGAA	4080
AGCGGGAATC	CAGCTGATCT	ATGGGATGGA	AGCCAATATC	GTGGAGGACC	GTGTCCTAT	4140
CGTCTATAAC	GAAGTGGAGA	TGGACTTGTC	AGAAGCAACC	TACGTGGTCT	TTGACGTGGA	4200
AACGACGGGA	CTTTCAAGCTA	TCTATAATGA	CTTGATTAG	GTTGCGGCTT	CTAAGATGTA	4260
CAAGGGGAAT	GTTATTGCTG	AATTGATGA	ATTATCAAT	CCTGGGCATC	CCTTGTCAAC	4320
CTTTACTACA	GAGTTAACTG	GAATTACAGA	TGATCATGTC	AAAAATGCCA	AACCACTAGA	4380
ACAAGTTTG	CAAGAACATT	AAAATTTG	CAAGGATACG	GTCCTAGTT	CCCACAATGC	4440
TACCTTGAC	GTTGGCTTTA	TGAATGCTAA	TTATGAGCGG	CATGATCTTC	CAAAGATTAG	4500
TCAGCCAGTT	ATTGATACGC	TGGAGTTTGC	TAGAAACCTC	TATCCTGAGT	ATAAACGCCA	4560
TGGTTTGGGG	CCTTGACCA	AGCGTTTGG	TGTGGCCTG	GAACATCACC	ACATGGCCAA	4620
CTACGATGCG	GAAGCGACTG	GTCGTCTGCT	TTTCATCTT	ATCAAAGAGG	TAGCAGAAAA	4680
ACATGGTGTG	ACCGATTAG	CTAGACTCAA	CATTGATCTA	ATCAGTCCAG	ATTCTTACAA	4740
AAAAGCTCGG	ATCAAGCATG	CGACCATCTA	TGTCAAGAAC	CAGGTAGGTC	AAAAAATAT	4800
CTTTAAGCTG	GTTTCCCTGT	CTAATACCAA	GTATTTGAA	GGAGTGCCAC	GGATTCCGAG	4860
AACGGTTCTA	GATGCCCATC	GAGAGGGCTT	GATTTTAGGT	TCAGCCTGTT	CAGAGGGTGA	4920

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AGTTTTGAC GTGGTCGTTT CTCAAGGTGT GGATGCGGCG GTTGAGGTGG CCAAGTATTA	4980
TGATTTATC GAGGTCATGC CACCGCTAT CTATGCACCC TTGATTGCCA AAGAGCAGGT	5040
CAAGGATATG GAGGAACCTCC AGACCATTAT CAAGAGTTG ATAGAGGTTG GAGACCCCT	5100
TGGCAAGCCT GTTCTGGCTA CGGGAAATGT TCACTATATC GAACCGGAAG AAGAGATTTA	5160
TCGTGAAATT ATCGTCCGTA GTTTGGGACA GGGTGCATG ATTAATCGAA CTATCGGTCA	5220
TGGTGAACAT GCCCAACCAAG CACCACTTCC AAAGGCTCAT TTTCGAACGA CTAATGAGAT	5280
GTTGGATGAA TTTGCCTTTT TGGGAGAGGA ACTGGCTCGT AAACTGGTTA TTGAAAACAC	5340
CAATGCCTTG GCAGAAAATAT TTGAATCCGT TGAAGTCGTT AAGGGTACT TGATAACGCC	5400
TTTCATCGAC AAGGCTGAAG AAACAGTTGC TGAGTTGACC TATAAGAAAG CTTTGAGAT	5460
TTATGAAAT CCGCTGCCAG ATATTGTTGA TTTGCGGATT GAAAAAAGAT TAACATCCAT	5520
ACTGGGAAT GGATTGCTG TGATTTATCT GGCACTCGAG ATGCTGGTGC AACGTTCTAA	5580
TGAACGGGGT TATTTGGTTG GTTCTCGTGG GTCTGTCGGA TCTAGTTCG TTGCGACCAT	5640
GATTGGGATT ACGGAGGTCA ATCCTCTCTC TCCTCACTAT GTCTGTGGTC AGTGTCAAGTA	5700
CAGTGAGTTT ATCACAGATG GTTCGTACGG TTCAGGATTG GATATGCCCA ATAAGGACTG	5760
TCCAAACTGT GGTCAACAAAC TCAGTAAAAA CGGACAGGAT ATTCCGTTTG AGACCTTCCT	5820
TGGTTTGAT GGGGATAAGG TTCCTGATAT TGACTTGAAC TTCTCGGGAG AAGATCAGCC	5880
TAGCGCCAC TTGGATGTGC GTGATATCTT TGGTGAAGAA TATGCCCTCC GTGCGGGAAC	5940
GGTTGGTACG GTAGCTGCCA AGACTGCCA TGGATTGTC AAAGGTACG AGCGAGATTA	6000
TGGCAAGTTT TATCGTGATG CAGAACTAGA ACGCCCTCGCT CAAGGAGCGG CGGGTGTCAA	6060
GGGGACAACCA GGCAACACCC CGGGGGGAAT CGTTGTTATT CCGAACTACA TGGATGTCTA	6120
CGATTTACG CCTGTCCAGT ATCCAGCAGA TGATGTCACG GCTGAATGGC AGACCACTCA	6180
CTTTAACTTC CACGATATCG ATGAGAACGT CCTCAAACTC GATGTAACGG GACATGATGA	6240
TCCGACTATG ATTCGAAACAC TTCAGGATTG GTCTGGTATT GACCCATAATA AAATTCCAT	6300
GGATGACGAA GGCCTGATGG CACTCTTTTC TGGGACTGAT GTGCTAGGGG TAACACCTGA	6360
ACAAATTGGA ACGCCTACGG GTATGTTGGG GATTCCAGAG TTTGGAACAA ATTCGTACG	6420
TGGAATGGTA GACGAAACCC ATCCGACAAC CTTTGCAGGA TTGCTTCAGC TGTCTGGCT	6480
GTCCCCACGGT ACTGATGTTT GGTGGGGAA TGCTCAGGAT CTGATTAAGC AAGGAATAGC	6540
GGACCTATCG ACTGTTATCG GTTGTGGGA CGACATCATG GTTACCTCA TGCATGCCGG	6600
TCTGGAACCT AAGATGGCCT TTACCAATTAT GGAACGGGTA CGTAAGGGTT TGTGGCTAAA	6660
GATTTCAGAA GAGGAGAGAA ATGGCTATAT CGAAGCAATG AAGGCTAATA AGGTGCCAGA	6720

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GTGGTATATC GAATCCTGTG GGAAAATTAA GTACATGTT CCTAAGGCC ATGCGGCAGC	6780
CTACGTTATG ATGGCCTTGC GTGTAGCTTA CTTCAAGGTT CACCATCCTA TTTATTACTA	6840
CTGTGCTTAC TTCTCCATTG GTGCTAAGGC TTTTGATATC AAGACCATGG GTGCGGGCTT	6900
GGAGGTCATC AAGCCAGAA TGGAAGAAAT CTCTGAAAAA CGGAAGAACAA ATGAAGCCTC	6960
TAATGTGGAA ATCGATCTCT ATACAACCTCT TGAGATTGTC AATGAGATGT GGGAACGAGG	7020
TTTCAAGTTT GGTAAATTAG ATCTCTACTG TAGTCAGGCG ACAGAGTTCC TCATCGACGG	7080
GGATACCCCTT ATCCCACCAT TTGTAGCAAT GGATGGTCTG GGAGAGAACG TTGCCAAGCA	7140
ACTGGTGGCGG GCGCGTGAAG AGGGAGAATT CCTCTCTAAA ACAGAACTAC GCAAGCGTGG	7200
TGGACTCTCA TCAACCTTGG TTGAAAAGAT GGATGAGATG GGTATTCTTG GAAATATGCC	7260
AGAGGATAAC CAGTTGAGTT TGTGTGATGA GTTGTTTAA AAAATTGCTT AATAATCTAT	7320
TAAAAGAGGC TAACGTATAT CCAATAGATT TACATTAGCT TTCTTTTTTG TTAAAATAGT	7380
CTATGGAAAG AGGGTGAGAG TATGTCAAAG ATGAGTATAA GCATCCGTCT GGATAGTGAG	7440
GTAAAGGAGC AGGCCAACAA GGTGTTTAGT AATCTGGAA TGGATATGAC AACAGCTATT	7500
AATATTTTCC TTCGTCAGGC AATTCAATAT CAGGGATTAC CTTTGATGT TAGACTAGAC	7560
GAAAATCGGA AGTTGCTCCA AGCGTTAACG GATTTAGACC AAAATCGTA TATGAGCCAG	7620
TCTTTGAAT CAGTCTCAGA TTTGATGGAG GACTTACGTG CTTAAGATTC GTTATCATAA	7680
ACAGTTAAA AAAGATTTA AGTTGGCTAT GAAGCGTGGT TTGAAGGCAG AATTATTAGA	7740
AGAAGTTTG AATTTCTGG TTCAAGAAAA AGAACATCCT GCCAGAACATC GTGATCATTC	7800
ATTGACGGCA TCCAAGCATT TTCAAGGAGT TCGTGAATGC CATAACCCAGC CAGATTGGCT	7860
TTTGGTTAT AAAGTAGACA AGTCGGAATT GATTTAAAT TTGCTGAGGA CAGGCAGTC	7920
CAGTGAATTA TTTAATCTA TTTAAGGGG GTTCTCATGA AACTAAGAAT ATTTGCGGAA	7980
GATAAGCCGG CTAAGAAGGT ATTTGAATAT CAATTAGAAC TTGCTGATCG TACAATTCTT	8040
CTATCGACAG CACTCTTGTCA AGGTGCTATT GCTTTAGCAG GAATCTTTTC TGCTTTGAAA	8100
GAAAATAAA AATAGAAAAG AGAAAACAGA ATGGTTTTAC CAAATTTAA AGAAAATCTA	8160
GAAAATATG CGAAATTGTT GGTTGCGAAC GGAATTAACG TGCAACCTGG TCACACTTG	8220
GCTCTCTCTA TTGATGTGGA GCAACGTGAA TTGGCACATC TAATCGTGAA AGAAGCTTAT	8280
GCCTTGGGTG CGCATGAGGT CATCGTTCAAG TGACAGATG ATGTGATTAA CCGTGAGAAA	8340
TTCCCTCCATG CCCCGATGGA GCGTTGGAC AATGTGCCAG AATACAAGAT TGCTGAGATG	8400
AACTATCTCT TGGAGAAATAA GGCTAGCCGT CTTGGAGTTC GTTCATCTGA TCCAGGTGCC	8460

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TTGAACGGAG TGGACGCTGA CAAGCTTC A GCTTCTGCTA AAGCTATGGG ACTTGCCATG	8520
AAGCCTATGC GTATCGAAC TCAACTAAC AAGGTTAGCT GGACTGTAGC AGCTGCAGCA	8580
GGACTTGGAGT GGGCTAAGAA AGCTTCCC A ATGCTGCGA GCGACGAAGA AGCAGTTGAT	8640
TTCCTTGGG ACCAAATTTC CAAAACTTGC CGTGTCTACG AAGCAGATCC TGTTAAGGCT	8700
TGGGAGGAAC ATGCAGCCAT TCTCAAGAGC AAGGCCGATA TGCTTAATAA GGAGCAATT	8760
TCAGCCCTTC ACTACACAGC GCCAGGAACA GATTTAACAC TTGGTTGCC AAAGAACAC	8820
GTTTGGGAAT CAGCTGGTGC TGTCATGCA CAGGGCGAAG AATTCTTGCC AAATATGCCA	8880
ACAGAAAGAGG TCTTCACAGC GCCTGACTTC CGTCGTGCAG ATGGTTATGT CACTTCTACA	8940
AAACCGCTTA GCTACAACGG AAATATCATT GAAGGCATTA AGGTGACCTT TAAGGATGGA	9000
CAAATCGTAG ATATCACTGC TGAGAAGGGT GATCAGGTTA TGAAAGACCT TGTCTTTGAA	9060
AATGGGGGTG CGCGTGCCTT GGGTGAATGT GCCTTGGTAC CAGATCCAAG TCCAATTCT	9120
CAGTCAGGCA TTACCTTCTT TAACACCCCTT TTCGATGAAA ATGCGTCAAA CCACCTGGCT	9180
ATCGGTGCAG CCTATGCGAC TAGCGTTGTT GATGGAGCGG AGATGAGCGA AGAGGAGCTT	9240
GAAGCTGCAG GGCTTAACCG TTCAGATGTT CACGTAGACT TTATGATTGG TTCTAACCAA	9300
ATGGATATCG ATGGTATTG TGAGGATGGA ACCGGGGTAC CTCTTTCCG TAATGGGAAT	9360
TGGGCAAATT AAGGAGATAA TATGTTAGGA AGTATGTTCG TTGGTCTCCT AGTGGGATT	9420
TTAGCAGGTG CTATGACCAA TCGTGGAGAG CGAATGGGAT GTTTTGAAA AATGTTCTC	9480
GGTTGGATCG GAGCCTTCT AGGTCACTTG CTCTTTGGAA CTTGGGGGCC AGTTTTATCA	9540
GGAACAGCTA TTATCCCAGC GATTTAGGA GCCATGATTG TTTTAGCTAT TTTTTGGAGA	9600
CGAGGAA	9607

(2) INFORMATION FOR SEQ ID NO: 81:

- (i) SEQUENCE CHARACTERISTICS:
- (A) LENGTH: 14231 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 81:

CTACAAAGATA ATTCCAGCTA TAACATCCGC TATAATAGTA AGAGCGAGCT CTATGATAAG	60
GCTCATTAGT TTCACCTCCT CTCACGAACC CATAGGAACG TAATCGGTAA CCGATGACAA	120
AAATAGTATA CCACAAATACA TTTAGATCAT CAAGGTCACT TAATTCTGA AATATCAGAT	180
CTAAGAGAAA AATCTTTAAA ATCAGAAAAA CGCATAATAT CAGGTGTGCA AAAACTTGAT	240

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ACTATGCGTT TTATTGTGGG AAGGTTTACT CCATTTCTC CTGAAATTGA GTTTTGTC	300
AGCCTCTGTT TTTAGGGTTG CTAAGAAAAT AATGTCATGT GGTGAATATT TGAAATCAG	360
TCAGCAGACA GAACGATACT CTTCGAAAAT CTCTCACAT CATGTCAGCT TCGCTTTCC	420
GTATATATGT GACTGACTTC ATCAGTTCTA TCTACAAACCT CAAAACAGTG TTTGAGCTG	480
ACTTGATCAA TTTCAAAATC TGACTTTGA GCAAGCTGAG ACTAGCTTCC TATTGATTT	540
TCATTGAATA TCAGAAACCC ATTCTCCATC AAATAATTG ACTGCGTCTA ATAATTGGT	600
ATCTGGCAGC GTGCTGAAA TAAAGGTTGT GSTATTTGGAG AGGGGATTAA TTTTAAAAAA	660
TCCAGTCTTG TAAAATTTAG AACTATCAAT CAGTAAGATG GTTTCATGGG CTTTGTCAAT	720
AATATTCTTT TTGAAATAG CTTGGCTGAG AGAACGTTCA TAAACATATT GGTCATCAAT	780
ACCTCTTGCT GAACAAAATG CTAATCGAT ATTAAAATGA TCTAATAAAAG AATTTCCCT	840
ATCATAGTTG ACCACGGAAC AGGATTGATG TTGACCTCG CCAGATGTGA TAAAGATTT	900
GGAGCTATCT TTAACAGTTT CAGATAGGGT TTGTCAGTA TGAAACCAT TTGAAAAAT	960
AATCAAATTA TCAAGTTCAAG AAAGATAGGG ACAGAGTTCG TAGACAGTAG TACTAGAAC	1020
TAGATAGATA CACATACCAG ACCGAATAAA GTCTTAGCG AGACTAGCGA TTGCTTTT	1080
TTGCCTAGTA CTTCTCCTT CACGTATTG ATGAGAAAGT TCAATTGTGT TCATAGAGGA	1140
CAGGTCACG TATCGTGCT TTCTTTGAT AAGACCTTGA TTTCTAAGA AAATTAAATC	1200
ACGACGTAAG GTACTTGTGC TCGAGAAAAGT GATTTCTGCC AGCTCTTTA CGGCAATTCT	1260
TTTTTCTTT TTGATAATT CAATCAATTCA AAGTACACGT TCATCTTTA TCATAAGCTC	1320
CTCCTAATT ATCATTCAA CTATATTATA GCACAAATTG GAGGAATTG AATTATTTT	1380
ATGAATATTG GGTAACACATT TGAACATTAT TCAAGTAAGC GTTCACATAT TGAAAAATA	1440
AAACGTGGGG ATTATAATAA AGTTAACGAGA GGACGAAGAG AGAAGAAAAA TGGAAAGCGGT	1500
TTTAGCAATA GATTTAGGTG CGACTTCTGG AAGAGCAATC GTGGTTACC TTTCTGAAA	1560
TAAACTAGTA ATGGAAGAAA TAAATCGCTT TTCTAATCTA CCTATTAGAG TAAAAGGGCA	1620
TTTATCTGG GATATTGACT TTCTACTAGC TAAAATTCTT GAAAGTATCC GCTTGGCTAA	1680
TACTAGTTAC AAGATTTAT CTATCGGTAT TGACACATGG GGAGTTGATT TTGGACTGAT	1740
TGATAATGAA GGTAAGCTGT TATTACAACC TGTTCATTAT CGTGATGAAA GAACAAAGGG	1800
AGTGTAAAG GAAATATCTG AAATGACTGA ATTAGAAAAA CTGTATTCAG AGACAGGAA	1860
TCAGATTATG GAGATAATAA CCTTGTCTCA ACTCTTTAAG GCACGTCAAG AATCTCCTGA	1920
CTCTTCTAT AAGACCAATA AGATTCTTT AATGCCAGAT TTGTTAATT ATCTCTTGAC	1980

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AGGTAAGTTT GCTACAGAAA AAAGCATTGC TTCAACAAC T CAATTATTTG ATCCCTAGGAG	2040
TCAAAATTGG AATCAGAATA TCTTAAAAC T ATTGAATTG GATTCATCTT TACTTCCTGA	2100
AATTGTTTCA GAGGGAAATG TTCTTGGAAAG GATAAAAGAG GAGTATGGTT TAGGCGATAT	2160
TCCTGTTGTG AATGTTGTG A GTCATGATAC AGCAAGCGCG ATTGTCAG TACCTAACAG	2220
AGAAGGTAGT TTATTTATTT CATCAGGTAC TTGGTCTTTG GTTGGAGTGG AACTTACTTC	2280
ACCGATTCTT ACTACCAGAAT CCTTCAGTTA TGGAATTACA AATGAAGTCG GTAAAGATGG	2340
AGTGATTACA TTTCTGAAGA ATTGTACAGG GTTGTGGATC ATAGAGGAAC TAAGACGTTC	2400
ATTTGAACGA AGAGGGAAAG CCTATTCTTT TGATGATATT AGGACAATGG TGGAGAAAGA	2460
AAAAGAAAAT CTTCCCTCTGA TTGATACTGA ATCAACTGAA TTTGCAACAG AATCTGATAT	2520
GCACAAGACT TTGACAGAAT ATCTAGCTT TCATCATGAA ACTAGAGAGT GGACAGATGG	2580
ACAACATATT AAGATTGTTT ATGAAAGCCT AGCTGAAACG TATAGGAAAG CGATAGAGTT	2640
ACTAGAAAGAA CTAACATCATA AGGTTTATAA GAGGATATAT GTGATTGGAG GAGGTGCTAG	2700
AGCCAGTTAC TTTAACCAAA TGATTGCTGA TAGAACTGGT AAAGAGGTTTC TTACAGGTTT	2760
GACTGAGGCT ACAGCTGTGG GGAATATTGT TGTCAGCTC ATAGCTATGG GACAATTAAA	2820
AGGGATGGAA GAGGCTCACC ATGTTATTGA GGAGTTCTA CAATTAGAGA GTTATTACTC	2880
CCAAAAGAAT TAAAAAGATT GAGAGTTGT AAATTGCTT CCCTCCCCCT TCTTAGCTTT	2940
TGTGCAGGAA GGGGGATAA TTGGTGAATT GAAAATATT TAGTGTGTTG ATATGAGGAG	3000
GACAAGGATG TCAGATGTAA AACAGAAATT AATTAAATAT GGTAAGAACG TAGTAGAAC	3060
AGATTTGACG AAAGAACAG GTGGAAATCT CAGCGTTTC GATCGTAAA AACATTGAT	3120
GGCAATTACC CCGTCGGGTA TTGATTCTT TGAAATCAA GAATCCGATA TTGTAGTGAT	3180
GGATATTAAT GGAAATGTTG TAGAGGGAGA ACGCTTGCCA TCTAGCGAAT GGTATATGCA	3240
TTTGATTCAA TATCAAACTC GTGATGATAT CGATGCAATT ATCCATGCTC ATACAACCTA	3300
TGCAACAGTA TTAGCTTGTC TCAGAGAACCC ACTTCCAGCG AGTCATTATA TGATTGCAGT	3360
GGCAGGGAAA GATGTTGGG TAGCTGAGTA TGCAACATAT GGCACGAAAG AATTGGCTGT	3420
GAATGCAGCT AAAGCAATGG AAGGTCGTAG AGCAGTTTA CTAGCGAATC ATGGAATT	3480
AGCAGGTGCA CAAAATTAT TGAATGCATT TAATATTGTT GAAGAAGTTG AATATTGTGC	3540
AAAAATTAT TGTAGCTA AGAATTGG AGAGCCAGTA GTTCTTCCTG ATGAGGAGAT	3600
GGAATTGATG GCAGAAAAAT TTAAACATA CGGTCAGAGA AAATAGGGAG GATATTAATG	3660
TTAAACATA TACCGAAAAA TATTCTCCA GATTATTGAGA AGACTTTAAT GGAAATGGGA	3720
CATGGAGATG AAATAGTATT AGCTGACGCG AATTATCCTT CTGCCTCATG TGCAAATAAG	3780

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CTAATTCGTT GTGATGGTGT AAATATTCCA GAATTATTAG ATTCCATTCT GTATTTAATG	3840
CCATTAGATA GTTACGTCGA TAGTTCAATT CAGTTATGA ACGTTGTTTC GGGTGATGAT	3900
ATTCCTAAGA TATGGGGTAC CTATAGACAG ATGATTGAAG GTCATGGTAC AGATCTTAAA	3960
ACGATTACTT ATCTTAGAAG AGAAGACTTT TATGAACGTA GTAAGAAAGC TTATGCTATT	4020
GTTGCTACAG GAGAAACTTC ACTTTATGCT AATATTATCC TTAAGAAAGG AGTAGTTGTT	4080
GAAAGAGAAA ATGTTCAATA GAGGAATTTT AGTTGCCAGT CATGGTAATT TTGCTAGCGG	4140
AGCTCTCATG ACCGCAGAAA TGTTTGTG TGAGACAACA AATGATAGAG TTAGGACATT	4200
AGGTTTGATG CCTGGAGAGA ATATTGTAGA GTTTGAGCAT TATTTTAAAA ATCAAGTGGA	4260
TGAACTGTTA GACTCAAATC AAGAGGTTAT CGTTTGACT GACTTGATTG GAGGAAGTCC	4320
TAATAATGTG GCTTGTAC GGTTTTTAAA TTTGGATTCA GTTGATATTG TAACAGGGTT	4380
TAATATCCCT CTCCCTAGTGG AATTAATATC AAGTTATGAT TCAAAAATCA ATTTAGAAGA	4440
AATTGTTCAC AATGCTAAA ATAGTTGTT TAATGTTAAA CAACAACCTTA ACGTAGAGGA	4500
GGAAGAAGAT TTATGCTAT AGAGTTGTT CGTATTGATG ACCGCTCTGGT ACATGGTCAA	4560
GTTGTCACTA CGTGGCTAAA AAAGTATGAT ATTGAGCAAG TTATCATTGT TAATGATCGC	4620
ATCTCAGAACG ATAAAACACG ACAATCTATT TAAAGATTT CTGCACCGGT AGGTTTAAAA	4680
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GGAGGTGTTAG CTCTAGGTGA AGAAGACAAA TATTATTTA AGAAAATGT TGATAAGGGA	4920
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TAAAATAAT TTAAGGGAGT ACAGTATATG CTATTCAACAC AACGATTACT GGTGACATTA	5040
GTTGGGATTA TTGCCACTAT TGACTATAAT GGACCGTTAT TTATGATTCA CCGTCCGTTA	5100
GTTACAAAGTG CAATGGTGG CTTAGTATTAA GGAGATTCA CCCAAGGTGT TCTTATTGGT	5160
TCAGCTCTTG AATTAACCTTG GCTCGGTGTA ACAGGTATTG GAGGTATAC TCCACCAAGAT	5220
ACTATTCAG GTGCGATTAT TGGTACTGCA TTTGGTATT TATCTGGTCA AGGAGAAACT	5280
GCTGGTATCG CTATAGCAGT TCCAATTGCA GTTGCTACCC AACAGTTGGA TGTTCTTGCA	5340
AAAACTTAG ATGTTTATTG TGTAAAAAA GCTGATAATG ATGCTAAAAA CGGAGATTAT	5400
TCAAAGATCG GTTTTATCA TTATTCAAGT TTGGTTTAA TCACGTTATT TAAAATTGTA	5460
CCAATTTCCTC TAGCTATTAT GCTTGGAGGG GAATATGTGG CAGACTTGTT TGCTAAGGTT	5520

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CCACCAATCG TTATCCAGGG ACTTAACCTCT GCAGGTGCTT TACTACCTTC AATTGGTTTT	5580
GGTATGCTTT TAAATATGAT GCTCAAGAAA AATATGTTGGG TATTCTTGTT GATTGGATTCTC	5640
ATTTGTTCTG TGATGGAGG AATGTCAACC ATTGGATCT CACTAGTTGG TATTGCGGTA	5700
GCATACTTCT ACAGATATGAT TGGAAAGCAAA CCACAAGAAA CAACTTCAG TAGTGATGTT	5760
GAGGAGGATC TTGATCTATG ATGAATAATA AAGTAACCAA AGTTGAACCTT AAAAAAGTTT	5820
TCAAACGAAG TTTTATGAT GGTTCTTCAT GGAACATATGA GAGAATGCAG AACCTAGGTT	5880
TTCTATATAC AATTCTTCCA GTATTGAAAA AACTATACCC AGACAAAGAT TCAGCTTCTC	5940
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GTATCTTTAT CGCCTTAATA TTGTTTAATA TTATTAATAT TCCTGTTAAA TATTCGGTT	6240
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ATGTGGCCT CTGCTGGAGG TAGAAAAGGT GTTCAGTTA AAGTAGTATG TCATTCTAAG	7260
GTTACGGATG GTGAAGTAAG TGTATTGGGA GAGACAATAG TTATTCGGAA TGCTACAGAG	7320

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GTATTTCTTT ATCTCAAATC AATGACGGAT TATTGGGAA ATATAGATAT TTCTTCTCTT	7380
CAGGGAGAAT TTAGTAGTAT TGATTACTTT ACAGAAAAAG ATGAACATGT AAAAAAATAT	7440
CAGGAGCAAT TTAATAGAGT TGATTTAAA CTAGACTATA GTAAAGGTTG TCTTAGCATT	7500
CCAACGAATC TACTTCTTGA AAACACTAAA AAGTATAGTA ACTACTTGAC TAACTTGTTA	7560
TTTCATTATG GAAGATATCT GTTAATATCG TCTAGTCAAC CGAATGGTT ACCTGCCAAT	7620
CTTCAAGGAA TATGGTGTGA TGAATTAAT CCAATTGGG GTTCTAAATA TACGATTAAT	7680
ATTAATACTC AAATGAATTA TTGGATGGTA GGTCCATGTG ATTTACCAGA ACTAGAATAT	7740
CCATTATTTG ATATGCTCGA AAGAATGAGA GAACCGGGAA GACTAACCGC TAAGAAAATG	7800
TATGGAGCTA GAGGTTTAC AGCACATCAT AATACGGATG GTTTTGGCGA TACGGCTCCC	7860
CAATCTCATG CCATGGGGGC TGCAATTGG GTATTAACTA TTCCATGGTT ATGTACTCAT	7920
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TTATGTTTG GATTGTTAGC AGGAGTTAAG GCTGATAATC GTGTTCAAAT GAGAACGACG	9060

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ATTAATAATG	AATGCCATT	GTTGCTTCT	CCGTTGTATG	GCAATGATAA	TGGTAACGGA	9120
TTATGGTGGG	GGAACACATT	GAAGGGAGCA	TGGGAAGCTA	TTCCCTGAAGA	TGTAAAGCCA	9180
TATGCAGCGA	TTGAACCTCA	TCCTGCAAAA	GTCTGTAAAC	CAACAAGTTG	TATTCACAGA	9240
GATACGAAAG	AATTGAGAGA	ATCGTATGTC	AAGATGTTGG	AGGAAGCTCA	AAGTCTAAAC	9300
ATTCCAGTTT	TCTTGGTTAT	TATGTCGGCT	GGAGAGCGTA	ATACAGTTCC	TCCAGAGTGG	9360
TTAGATGAAC	AATCCAAAAA	GTATAGTGTG	TTAAAAGGTG	TTTTAAATAT	TGAGAATTAT	9420
TGGATTTACA	ATAACCAGTT	AGCTCCGCAT	AGTGCTAAAT	ATTTGGAAGT	TTGTGCCAAA	9480
TATGGAGCGC	ATTTTATCTG	GCATGATCAT	GAAAAATGGT	TCTGGAAAC	TATTATGAAT	9540
GATCCGACAT	TCTTGAAAGC	GACTCAAAAA	TATCATAAAA	ATTTGGTGTGTT	GGCAACTAAA	9600
AATACGCCAA	TAAGAGATGA	TGCGGGTACA	GATTCTATCG	TTAGTGGATT	TTGGTTGAGT	9660
GGCTTATGTG	ATAACTGGGG	CTCATCAACA	GATACATGGA	AATGGTGGGA	AAAACATTAT	9720
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TCAATGATTG	CTATGGAAAT	GATGAATGTA	TATACTGGGG	GAGGCACAGT	TTATAATTTC	9840
GAATGTGCCG	CGTATACATT	TATGACAAAT	GATGTACCAA	CTCCAGCATT	TACTAAAGGT	9900
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CTTTATCCAA	AACTTTATGA	AGGAGATGGG	TATGCTCAGC	GTGTAGGTAA	TTCTGGTAT	10260
ATTTATAATA	GTAATGCTAA	TATCAATAAA	AATCAGCAAG	TAATGTTGCC	TATGTATACT	10320
AATAATACAA	AGTCGTTATC	GTTAGATTG	ACGCCACATA	CTTACGCTGT	TGTTAAAGAA	10380
AATCCAAATA	ATTTACATAT	TTTATTGAAT	AATTACAGGA	CAGATAAGAC	AGCTATGTGG	10440
GCATTATCAG	GAATTTTGA	TGCATCAAA	AGTTGGAAGA	AAGAAGAATT	AGAGTTAGCG	10500
AACTGGATAA	GCAAAAATTA	TTCCATCAAT	CCTGTAGATA	ATGACTTTAG	GACAACAACA	10560
CTTACATTAA	AAGGGCATAAC	TGGTCATAAA	CCTCAGATAAA	ATATAAGTGG	CGATAAAAAT	10620
CATTATACTT	ATACAGAAA	TTGGGATGAG	AATACCCATG	TTTATACCAT	TACGGTTAAT	10680
CATAATGGAA	TGGTAGAGAT	GTCTATAAAAT	ACTGAGGGGA	CAGGTCCAGT	CTCTTTCCCCA	10740
ACACCAAGATA	AATTTAATGA	TGGTAATTG	AATATAGCAT	ATGCAAAACC	AACAACACAA	10800
AGTTCTGTAG	ATTACAATGG	AGACCCATAAT	AGAGCTGTGG	ATGGTAACAG	AAATGGTAAT	10860

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TTTAACCTCTG GTTCGGTAAC ACACACTAGG GCAGATAATC CCTCTTGGTG GGAAGTCGAT 10920
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 CAACGCTCTAT CTAATTTGA TGTGATTCTA TATGACAATA ATAGAAACGA AGTTGCTAAG 11040
 AACACATGTTA ATAATTTGTC GGGTGAATCT CTTAGTCTAG ATTTCAAAGA AAAAGGAGCA 11100
 AGGTATATTA AAGTTAAATT ACTAACCGAGT GGAGTGCCTT TGAGTTTAGC AGAAGTAGAG 11160
 GTTTTAGAG AATCAGATGG TAAGCAATCT GAAGAGGATA TAGATAAAAT AACAGAAGAT 11220
 AAAGTAGTCT CTACAAATAA GGTAGCTACT CAAAGTTCAA CCAATTATGA GGGTGTAGCT 11280
 GCTTTAGCAG TTGATGGTA TAAAGATGGA GATTACGGAC ATCATTCCGT GACTCATACT 11340
 AAGGCAGATT CTAACGCTTG GTGGCAGGTC GATCTGGGAG AAGAGTTTAC GGTTTCTAAA 11400
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 GCAGCTATTC CGTTAAGTTT AGCGGAAGTT GAAAGTCTATG GTTCAAAGAG AACTCCGAAG 11640
 AAACTTCTA ATATTGCATT AACAAAAGAA ACTCGACAGA GTTCAACGGA TTACAATGGT 11700
 TTTTCTCGTC TAGCAGTTGA TGGAAATAAA AACGGAGATT ATGGTCATCA TTCAGTGACT 11760
 CATAACAAAG AAGATTCTCC TTCATGGTGG GAGATAGATT TAGCACAAAC CGAAGAATT 11820
 GAAAAGTTAA TTATTTATAA TAGAACAGAT GCTGAAATTC AGAGATTATC AAATTTTGAT 11880
 ATTATTATAT ATGATTCAAA TGATTATGAA CTTTTTACAC AACATATTGA CAGTTAGAA 11940
 AGCAATAATC TATCCATAGA CTTAAAAGGA CTGAAGGGAA AAAAGGTTAG AATTCTTTG 12000
 AGAAGCGCAG GAATCCCTTT AAGTTTAGCA GAGGTAGAGG TTTATCTTA TAAGTAATTT 12060
 TAAAATTAT CACCCAGGCT ACCGTTAAATA TAATGGAGAT GGTAGTATGA AAGAAACAGA 12120
 AAAATAAGAG GAAAATAGTA TGATTCAACA TCCACGTATT GGGATTCGTC CGACTATTGA 12180
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 TGTGGCAGAT TTGATTTCAA GCACATTGAA ATATCCAGAT GGGGAACCTG TGGAATGCC 12300
 GATTTCTCCA TCTACTATTG GCCGTGTACC AGAGGCTGCA GCTTCCCATG AGTTGTTAA 12360
 AAAATCAAAT GTTGCGCAA CAATTACAGT TACACCATGC TGGTGTATG GTAGTGAAC 12420
 TATGGATATG TCTCCAGATA TTCTCATGC TATTTGGGA TTTAATGGGA CAGAACGCC 12480
 AGGAGCTGTC TATCTTGCAG CTGTACTAGC TTCACATGCT CAAAAAGGGAA TTCCAGCCTT 12540
 TGGGATTTAT GGAAGAGATG TTCAGGAAGC TAGTGACACA GATATTCCAG AAGATGTCAA 12600

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AGAAAAACTT TTACGCTATG CGCGTGCAGC TCTTGCAACT GGCTTGATGA GAGACACTGC	12660
TTACCTATCA ATGGGTAGTG TTTCGATGGG GATTGGTGGT TCTATTGTAATCCGGATT	12720
CTTCCAAGAA TACTTAGGAA TGCGAAATGA ATCGGTAGAT ATGACGGAGT TCACCGGCCG	12780
TATGGACCGT GGTATTTACG ACCCTGAAGA GTTCAACGT GCGCTCAAAT GGGTAAAGA	12840
AAACGTAAGAA GAAGGATTG ACCATAACCG TGAAGACCTT GTTTAAGCC GTGAAGAAAA	12900
AGATAGACAA TGGGATTTG TTATTAAGAT GTTCATGATT GGACGTGACT TAATGGTTGG	12960
TAACCCAAGA CTTGCTGAAC TTGGTTTGA GGAAGAACG GTTGGTCACC ATGCTTTAGT	13020
AGCTGGTTTC CAAGGTCAAC GTCAGTGGAC AGACCATTTC CCAAATGGGG ACTTTATGGA	13080
AACTTTCTC AATACTCAGT TTGACTGGAA TGGTATTGAA AAACCATTTG TATTTGCGAC	13140
AGAGAAATGAT TCACTAAATG GTGTGTCTAT GCTCTTAAAT TATCTATTAA CAAATACTCC	13200
ACAAATCTT GCTGATGTGC GTACTTATG GAGCCAGAG GCTGTTAAAC GTGTAACGGG	13260
ACATACTTTA GAGGGTCGTG CTGCAGCTGG CTTCTTACAT CTAATCAACT CTGGTTCTTG	13320
TACATTGGAT GGTACAGGTC AAGCTACTCG AGATGGCAA CCTATTATGA AACCAATTCTG	13380
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CCCGGAATAC TTCCGTGGAG GAGGATTCTC AACTCGTTTC TTGACGAAGG GGGATATGCC	13500
AGTAACAATG GTACGCTCA ATCTTCTAAA AGGGGTTGGT CCAGTGCCTAC AAATTGCAGA	13560
AGGTTACACA CTTGAACTTC CTGAAGATGT TCACCATACT TTGATAATC GTACAGATCC	13620
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CTATGACGTC ATGAAATAATT GGGGAGCTAA TCACGGAGCC ATAACATATG GACACATTGG	13740
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TGAGGAAGAT ATCTTAGAC CTAAGGAAATTG GTCCCTTATTT GGAACAGAAG ATCTAGAAC	13860
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ATTGAAAACG AATACAAAAA GTAATATAAT GATGTTAAAT AGATAGCGCG GAGGCGCAGG	14040
AGGAAAATTA TATGGCTATA TTTTATGTT CGGCAGTCAA CCTTATTGGA AAAGGTGTTG	14100
TAAATGAAGT GGGTCCTTAT ATCAAGGAAC TTGGCTATAA AAAGGCACCTT TTGGTGACAG	14160
ATAAGTACAT CGAAGGCAGT GATATTTAC CTAAGACTTT AAAACCACTG GATACAGAAG	14220
GAATCGAATA T	14231

(2) INFORMATION FOR SEQ ID NO: 82:

(i) SEQUENCE CHARACTERISTICS:

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- (A) LENGTH: 16995 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 82:

AGTTCTCTTA ACTTTTTAG GATGGCATTC TCCGCTCTCA GGTACTCATT TTCTGCTgAA	60
GACGTTCTAA TTCTGTCCTC TCTTCAGGTC TCGTTTTGG CTTACGTCCC ATTTTAGGTA	120
CTCTCCCTCT TGTTTCTCA ACAATAGTAT ACCCGTTTT CCTGTATTGT GCTAGGCCAGT	180
TAAGAAGTAT CGTACCACTT GGGAGACCGT ATTCAAGAGA AACTCTATCT TTAGTCCAGC	240
CTTCATGTCA GACTTTATTAA CTCATTTCTT GTTTAAATC AGGAGAATAG TAACGATTTT	300
TTCCCTTTTT GACGAACCT ATTCCGTAAC GATCAATCAA TTTAACATG TACCTAATAT	360
TAGAATTGCT TATCCAAAT TTATTTGAAA GCTTCTCTAA GCTATATCCT TGTTTCTAA	420
GTTCATAGAT CTGAACTTTA TCATCATAAG TTAGTTTCAT AATAAAAACA CCCCAAAAGT	480
TAGATTTTT CTGCTAACT TTGGGGGTGT AGTTCATGTA CACCTGATAT GATGCGTTTT	540
ATAATTTTA AGCCTTTTG CCCAGCCTCG TCAAAAGTAA TGTTTGACA CAAAATCTGT	600
GACAAAACMT TAGTTTAAAC GGTTTTAAC TTGTATATA CTAGTTTAA GAAAAGGAGG	660
ATGATCTAAT GGAAGAAAAA GTATCATTGA AAGTCAGGGT TCAAAACTA GGGACATCGC	720
TTTCNNTAT CGTTATGCCA AATATTGGAG CATTTATTGC TTGGGGAGTA TTGACTGCC	780
TCTTTATCGC TGATGGCTAT CTGCCAAATG AACAGTTAGC TACTGTTGTT GGTCCTATGT	840
TAACGTATTT ATTGCCAATC CTGATTGGTT ACACAGGTGG ATATATGATC CATGGCCAAC	900
GTGGTGCCTG TGTAGGAGCT ATTGCTACTG TTGGTCAAT CACAGGTTCT AGTGTTCCTA	960
TGTTTATCGG AGCTATGGTA ATGGGCCAC TGGGAGGATG GACTATCAAG AAATTGATG	1020
AGAAGTCCA GGAAAAAATT CGTCCCGGAT TTGAAATGTT AGTTAATAAC TTCTCAGCTG	1080
GTCTCGTTGG TTTGCTTAA TTGCTTTGG CTTCTACGC AATCGGCTCA GTCGTATCGA	1140
CTCTTACTGG AGCTCTTGGG AATGGTGTG AGGCTATTGT CAATGCTCGC CTCCCTCCTA	1200
TGGCTAATAT TATCATCGAA CCGGCTAAAG TCCTTTCCCT CAATAATGCC CTCAATCATG	1260
GCATTTTAC TCCTCTGGGA GTAGAACAGG TAGCTCAAGC TGGTAAGTCA ATTCTCTTCC	1320
TATTGGAAGC TAATCCTGGA CCAGGTCTGG GAATTCTATT AGCTTATGCT GTATTGCGTA	1380
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ATGAAATTAA CTTCCTTAT GTTATGATGA AGCCTACTCT ATTTTAGCT GCTATGGCAG	1500

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GAGGTATCTC TGGAACTTTT ACTTTCAAC TCTTAGAGCG TGGCTTAAA TCTCCAGCTT	1560
CACCAAGGTTTC TATTATTGCG ATTATAGCTA CGGCCCAAA AGGTGTTGG CCCCCATCTAA	1620
ATGTTCTTTT AGGTGTTTA GTGGCAGCAG TTGTTCTTT CCTGTAGCA GCCCTTATTC	1680
TTCATGCAGA CAAGTCAACT GAGGATTCGC TCGAAGCTGC TCAGGCGGCT ACCCAAGCAG	1740
CTAAGGCTCA GTCTAAAGGT CAGTTAGTAT CAACTCTGT TGATGCAGTT GTTCGACAG	1800
ACTCAGTGA AAAAACATT TTCGCCTGCG ATGCTGGTAT GGGAAAGCTCT GCTATGGGAG	1860
CTAGTATTCT TCGAGATAAG GTTAAAAAAG CAGGTCTAGA GATTCCAGTA TCTAACAGG	1920
CAATCTCAA TTTGCTTGAT ACACCAAAAA CATTAATTGT TACTCAGGAA GAACTGACAC	1980
CAAGAGCTAA AGACAAGAGT CCAAGTGCTA TTCATGTTTC TGTTGATAAT TTCTTAGCGT	2040
CCTCTCGTTA TGATGAAATT GTAGCTTCAT TAACAGGAGC TTCTCCAATA GCAGAAATTG	2100
AAGGAGATAT ACCAACTTCA GCACCAAGTAG ATAGTCAGGA AAGTGACCTT AACCATATTG	2160
ATGCTGTAGT AGTTGCTTAT GGTAAGCAC AGGGAACTGC AACTATGGGC TGTGAAACGA	2220
TTCGGGCTAT TTTTAGAAC AAGAATATTG GTATTCCAGT TTCTACTGCC AAAATTCA	2280
AATTAGGTGA ATTTAATTCT AAAAACATAA TGATTGTAAC AACTATTCTT TTACAGGCAG	2340
AAGTGCAGCA AGCAGCACCG AATTCTCAAT TTCTTATTGT GGATAGTTTA GTAACAACAC	2400
CAGAATATGA CAAAATGGCT GCTAGAATGT ACAAAATAGAA CTAGAGGTTT CTAATTACG	2460
AATGCTTATA ACCAAACGAG AAGAACAAATT ATTGAAGGCT TTCCCTACATG TAGGGAAAGCT	2520
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TATCTCAGTA GCAGATTTT CAACCGTAA TTTTGGGAGC TTGATATTAA TAGAACAGA	3000
TAGAACTGGG CTGGCCAGTC AGATTGTTAA TAAGCAACTG TCAGGTTTC CAGATATGGA	3060
TGCTAGGATG AAGATGTTT TTGCGATCTT GTTATCTCTT ATAGGTCAAG AGCAAAACAT	3120
TGAAAATTCA CCTAATACTA GTAAGCAGGC TTGGAAATT TCTCAAAAAA TTTTCAAGC	3180
TTACTCTAAG CAGACTGCAC AATTTATAG TATTCAGGAA ATTATCTATT TTGCGAGCAT	3240
CTTGGATGAA TTAATCATTAA AACGTCAGGA CAATCCGCTC TTTACGGAGA AATTTGATGG	3300

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TGAATTTTC TACAATATTT CAAATCTGAT TGATACGGTT TCCATGTATA CCAAGATTGA	3360
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CGTCCCTATC CTTCAGG GTGAAAATTT GCCAGAATCT ATCCAGATTT TAGTTGAAAG	3480
GAATAAAATTT CTTTATACAG TCATCAGTCT TTTAGTGAAT GATATTTTC CGAAATATCT	3540
TCATACAGAG TATGAGTATG GCATGATTGC CCTACATTTT ATCTCTAGCT TAGGCCGTAG	3600
TCCAGAGATT TATCCAGTCC GTGTTTGCT TTTAACGGAT GAACGTCGGG TCACTAGAGA	3660
TTTATTAGTC AGTAAAATTA AGAGTGTGTC TCCTTTGTA GAGTTGATAG ATATTCAATC	3720
TCTAGTAGAT TACCACAGTA TTGATCTCAG TCAGTATGAT TATATTTTAT CTACCAAGCC	3780
GCTGACTAAT CAGGAAATCG ATGTAATTTC TAGTTTCCA ACCGTCAAAG AATTGCTTGA	3840
ATTACAGGAA CGACTTCAGT ATGTACAGGC ACATCGTACA ATTGTCGCGC GTGATGCTAT	3900
CGCTCCAGAG AAAAGTTATG ACTTCCAAGA TTATTTAATA TCTAGTAGTC AGCTTTGAG	3960
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CATCCAATAT CAGAGAATG TGAGTGACAG AGCTTACCTA ACAAGAAAAT TGTTATCTCA	4080
CTTCCAGAAT AGTCCTATGG CTATTCCCAA TACTGGTCTG GTGCTTTAC ATAGTCAGTC	4140
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AGCTAGCGAG GAAGCTAGAG ATTTAATGAC AGCTATTAGT CAGTCGATTA TTGAAAATCA	4320
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TATTTTTAAC GAAAAAATTA AGAAATTGGA GAACTAATAT GAAACTTGAA AAACATTTGA	4440
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AAACATTCTG TTCAATTTGG TGCCGGTAAT ATCGTCGCGT GTTTATAGG TGAAATTCTA	4980
TTTAAAAATG GTTCCATAT TGATTTGTTG GATGTCAATA ATCAGATAAT TCATGCTCTG	5040

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AATGAAAAGG GCAAGTATGA AATTGAAATT GCACAGAAAG GACAGTCTCG TATAGAAGTA	5100
ACTAATGTGG CTGGCATTAA TAGCAAAGAA CATCCTGAGC AAGTCATTGA AGCGATTCAA	5160
AAGACGGATA TTATTACTAC TGCAATCGGA CCTAATATAC TCCCTTTAT CGCCGAACCT	5220
CTAGCCAAAG GAATCGAACG TCGCCGAGTT GCAGGAAATA CACAGGCATT GGATGTTATG	5280
GCCTGTGAAA ATATGATTGG CGGGTCTCAA TTTCTTTATC AAGAAGTCAA GAAATATTTA	5340
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TATGAAGAAG ATTTAGAACCC CTTTATTGAG CGAAAACCTT TTTCAAGTCAA TTCTGGACAT	5580
GCAACTTCAG CTTACATTGG TGCGCATTAT GGTGCCAAGA CAATTTGGAG AGCTCTTCAT	5640
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GAAGTAGAGG TGTACTATTTC TAGTTCAAT CTACTATATA ACTGAAAAAT TAGATAAATT	6660
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TGAGTGTGTTT AGTTAGGAAA AAGGCTTGTT GTCTATAATT GTCTGCATTA GTCTAGATT	6780
TATTTATAGA AAATGTTATA ATAGACTGTA TTTAAAAAAT TTTAAGGAGA AATGACAGAA	6840

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AAGCTCTTA TCAAGATGCA ATGAACGCAC TTTGCCAAA CGCTTATGAA GCAGCTGTAA	7080
AAGAACGCTGG TCTTGAAAGTG GTTGCCCAAC CAAAAATTGA CGTAACTTCA ATGGAAAAAG	7140
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GGGGGTAAGA GACTTCATTA CCATCAAAAT CAATGACAAT CTCGTCTTGT TTGATTTCGG	8640
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CATACTCAAT TCCCAGTTCT TGCTCGACAG TAGTGATATG TTTTGAAATG GTTCTAAAT	9600
CATGACTCTT CTGTTTCTT TTTCAGGA TACGAACCAA GTGACTGCGG ATGCCCTTCT	9660
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CGATCAGGTA GGAGAGCTCT TGGCAACTT GGCTGGGTC TAGTTCCACG GGACGGGAAG	9780
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CGCCATAGGT ATCCACGATT TTCTGAGCTG TCTTGAGACCC AATCCCTTG AAATGGCTAC	10260
TTGAAAAGTA CTTGACCAAG CCCTTACTAG TTGGTTTGC GCGATCATAA CGACTGATTT	10320
GCAGTTGTTGTC TCCATACTTG GAGTGTGGA CAATTGCCC CCAAAAGTA TAGTCTTCGC	10380

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CCTCAATTAC ATCAGCCATG GTTCCGTGTA CAATGATTTC AAAATCATCA AAATCCTCTG	10440
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TAATCCGTTC AATAGTTCTC GAAAATAAA CTTCCATAAA ATTCCCTTGCA ATGAATAGGT	10560
GAGAGTTGGG ATTGTTTTA TTTTATACTC TTCGAAATAA TCTTCAAACC ACgtCAGCTT	10620
CCATCTGCAA CCTCAAAACA GTATTTGAG CTGACTTCGT CAGTCTATC CACAACCTCA	10680
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GTAGTTGACA TCAACTGTGA AGGCTTGAGC TTTTGAGCG ATACTCTAA AGAAAGTTCC	11040
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TTTGATATAG TCTGCTAGAT AAGGCTCGTC CGTTTCTTG TCATTGATGT AGAGTTTATC	11160
ATTTTCGTAAC CGAATGGTGT CGCCAGGCAT TCCAATCAG CGCTTGACGA TGTCCCTTATT	11220
GCCATCTTCC TCAATGGGCCA CCACGATATC AAAACGGTCA ATAGGAAGGT GTTTTACAAC	11280
GAAGAGAAATT TCGCCATCGG CTAGGGTCGG ATCCATGGAA TGTCCCTCTA CGCGAACATT	11340
GCTCCAAAAA AAGATACGAC TTAAAGCTAG TAATGACAGA ATTAGGAGGA ACAATCCCCA	11400
CTCTTTTAAG AAATTTTAA ATGAATTCA AACTTACCTT TCTAAGCGTT TTTTCGCTTT	11460
TTCAGTGTGTT TTAAAGTGCAT ATTTGGCGCA GAAGCTGAGT CCCTGCATAC CATAGGCTTG	11520
CAAATCTGG CTAGCCACCT TGTCAAGAAGC CGTTCCAGCT CCACCTGGGA GCTGATAACC	11580
CAGTTCTCGT CCCAAATTTT CAAGATTTTC CAGAAAGAGA TCACCGCAA TGACAGAAGA	11640
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AGCCAGACTG GTTTGATAGT GTTCAAGAAA AGCCTGAATA TCCTTTTCGC TTGGTGTGAG	12300
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AAAATTAGCG AATTTGGTA TAATATCGTG AGGTGAATT TATGGCAAAT CTAATCGAT	12420
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GCGCAGATGA TGAAACAATC GCTCTTTGT TGCCAGTCAA CTGTTTATCA ACTCAGCTCA	12600
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AGAGCAATTC CATTATTCG TCCCTTATGC AAATTCGAG GAAGGTCAGG GGACTTTCTT	12900
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ATCTGGACTT	GATTCGTGCCT	AAGGTTCGAT	TTATCCAAGA	AAGACAAGCA	GTCGTCGCCTC	14220
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AAGAAGCCAT	GAATGAGCTA	GATACTTCA	TCGACCAAGC	CTTGCTTAAC	AAATGGCTC	15540
AAGTTGATAT	CATCCATGGT	ATCGGAACAG	GAGTCATCCG	TGAAGGAGTT	ACCAAATACT	15600
TGCAAAGAAA	CAAACATGTC	AAGAGTTTCG	GCTATGCC	ACAAAATGCT	GGAGGCAGTG	15660

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GTGCGACTAT	TGTCACTTT	AAAGGATAGC	AGTATTCTGG	ACTTTATAAA	GTAAAAACTG	15720
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AACTGCTCAA	ACTCTCAGGT	AAAAGGACAG	AGCTAGGATA	GACCGCTTT	TAGCATTAT	15960
CTAACGCATTC	CAGAGTACAT	GTATCTTGCA	TGTGCTCTTT	CTTTGGGGT	TGAAACGATA	16020
GGAGAGGAA	ATGTTAGAAT	TGCTTAAATC	AATCGATGCT	TTTGCTTGGG	GACCGCCCCCT	16080
CTTGATTTA	TTGGTCGAA	CAGGGATTTA	CCTAACTATT	CCGCTAGGAC	TCTTGCAGGT	16140
TTTGCCTCTA	CCCAAGGCCT	TTCAGCTTAT	TTTTATCCAG	GATAAGGGAC	ATGGTGTGAT	16200
ATCCAGTTTT	GCAGCTCTGT	GTACAGCCTT	GGCATCAACT	GTTGGAACAG	GAAATATCAT	16260
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CAAGGACGAC	CATGGTGCAG	TAGCGGGAGG	TCCCATGCAT	TATATCCTTC	TAGGGATGGG	16440
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GATTTCGCCA	GCCATCACAG	CTCTCGTCTT	GTCTGTCTTT	GTAGCGATTG	CAGTCTTGG	16620
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TTTAGCTCTT	ACCTCAGCTT	TTAGTCCCTT	TGCTCCGGTA	GGTGGATTG	CTGGTGCTAG	16800
CGTTTCGGATG	GCTATTCAA	ATGGTGTGGC	GGCTGGTGTG	TTCTCAAACG	AATCTGGTCT	16860
GGGTTCTGCT	CCTATTGCA	CTGCAGCTGC	CAAGACAAAT	GAACCAAGTAG	AGCAAGGTTT	16920
GATTTCCATG	ACAGGAACCT	TTATTGATAC	CCTCATCATT	TGTACTCTAA	CTGGTTTGAC	16980
CATCTTGGTA	ACTGG					16995

(2) INFORMATION FOR SEQ ID NO: 83:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 28473 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 83:

CCGGGGCTTT	TGTAGTATAA	TAGAGATACG	TTTTGAAAGT	AGGAGGTATC	TATGGACTTA	60
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ACTAAGCGCT TTAATAAAACA GTTAGATAAAA ATTCAAGTTT CGTTGATTG TCAGTTTGAC	120
CAGGCTATTT CGGAGATTCC TGGGGTCTTG CGTTTGACCT TGGGGGAACC TGATTTACA	180
ACGCCAGACC ATGTCAGGA GGCGGGCAAG CGAGCGATTG ATCAGAACCA ATCCTACTAT	240
ACAGGGATGA GTGGTCTGCT GACTCTACGT CAGGCAGCCA GTGACTTTGT TAAGGAAAAG	300
TACCAAATGG ACTATGCTCC TGAAAATGAA ATCTTGGTTA CAATTGGGGC GACAGAGGCT	360
TTATCTGCGA CTTTGACGGC TATTTGGAA GAGGGAGACA AGGTACTTTT GCCAGCTCCT	420
GCTTATCCAG GCTATGAACC GATTGTTAAC TTAGTTGGG CAGAAATTGT TGAGATTGAT	480
ACGACTGAAA ATGGTTTGT CTTGACTCCT GAGATGTTGG AGAAGGCCAT TTTGGAGCAG	540
GGTGATAAGC TCAAGGCGGT TATTCTCAAC TATCCAGCCA ATCCGACAGG AATTACCTAC	600
AGTCGAGAGC AGTTAGAGGC CTTGGCAGCT GTTTTACGCA AGTACGAAAT TTTTGTGTC	660
TGTGATGAGG TTTACTCAGA ATTGACCTAC ACAGGCGAAG CCATGTTGCT CTAGGAACGA	720
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CTGGTAAAAA CGATGCGGAC CCATGAAGAA GGAATATATC CAACGTCGGG ACTATATCAT	960
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GCACGATTTTC TCTTGCGAAT CAATGATGAC GGACTCAGTT ACATCGAAGA CTATCATGAG	1440
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GCAGCTCTTG CAGATGCTAG TTTCGAGGAC AATCAGCAGG ATGCTCCCTT GTTTGCTTTT	1560
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TGCCATCGGG TTGGTCAGGC TTTTGACTTT TCTTTCAAAT ATGGAGCCTG CCTCTGTCCA	1740
GAGCATTATC ATGAGGATAA GAGACGTTGT CATCTCAATC CCAATATCCC CTATCTGCTC	1800

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						3600

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ATCATTGGAA AGAAAATAAG CTGGTTAAGA AGGGGGATCT TTGGTTCAA TACCAAGAAG	7140

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CAAGACTGCG ACTGGTGGCT ACGATGGTCA TGGACAAAAG GTTATTGCTT CAGAAGCAGA	21240
CTTGGAAAGCA GCCTATGCGC TAGCAGACTC AGCAGACTGC GTCTTGGAAAG AATTGTCAA	21300

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CTTTGACCTT GAGATTCTG TCATCGTGTG AGGAATGGC AAGGAGGTGA CGTTTTCCC 21360
 AGTTCAAGAA AATATCCACC GCAACAATAT CCTGTCTAAG ACCATCGTAC CAGCCCGCAT 21420
 TTCTGAAAGT CTAGTAGACA AGGCTAAAGC TATGGCAGTG CGAATCGCAG AACAACTCAA 21480
 CTTGTCTGGA ACTCTCTGTG TGAAAATGTT TGCGACAGCT GATGACATCA TTGTCAATGA 21540
 AATCGCCCCA CGACCACATA ACTCTGGGCA CTATTCTATT GAAGCCTGTG ATTTCTCTCA 21600
 GTTTGACACC CATATTCTGG GTGTTCTCGG AGCACCATTA CCAGTCATCA AACTCCATGC 21660
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 GTTTGAGTAG AAATCTTGGT TTACCTAGAT AGCTTATTCC CAACAGCTTA AGAAGAAAAGG 22140
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 GTTTCTGAGA CTCTTGGTGA AGAGCGCAAG TGGGTTCACT ATGGGTTAAC TTCTACTGAC 22440
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CAGAAGTAAC ATCACGTCTC ACACAAGAAG AAATCGATGA AATCTTCAAC CCAGTTTATT	23400
ACACAAACG AGTGGATGAT ATCTTGAAAC GTCTGGACT AGGTGATTAA TTAAAAAAATA	23460
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TCGTGAGTTC CTGTTTCAGG AAGTTTTTC TCTGTTACCA CAGGAGCTGG ATCTTGAGGA	23640
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GCAAGACAC TTTCATCAC CTTGTGAGTT ACAGGAGCGC CAACTTCAAC CACTTGGTT	24060
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GAATGAGTTA CAGCTGGTTT GAGGCCCTGA AGAGCGGCTT TTAGGTTGGC TACAAGCGTG	24360
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AGACCGATAC CTTTATCTTT GGCATAGTTA ATCAGATCTG TCATTTGACT TTCTGTTAAC	26640
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TCGTCACTGG CATAGGTCTT GCCGTTAGCT GTGATGCTCA TATCGTCCAA CATGAAACGG	26760
AGTCCATCAT TTCCGACTAA TAGGTGTAAA TCAGTGTAGC CATAATGTTT CGCTTTATCG	26820
ATGATTTCCT TGAGCTGTTc TGGTGAGAAA TATTTACGTC CAGCATCAAT AGAAACAATT	26880
TTCTTTTCG CTAGTTTTC ATTACAGTT GCAGCACGTT CCTTCCTGC CTCTGTTGCC	26940
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AGCTTGTGTT TGCGTGCACG TGAAACGAAG GTTCTTTTC CTTGTTGGCG GACAATATAG	28320
CCATCTTGG CAAGGTCGTT TAAGGCGCGA ACAACTGTGA TAGAGCTGAC ATCGTACATT	28380

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GAAATGAGTT CTGCTTCAGT GTAAAATTAA TCTCCACTGC TAAACTGCC AGAGATGATT	28440
TTATTTTTA ATTGGTCTTT TATGTATTGA TGG	28473

(2) INFORMATION FOR SEQ ID NO: 84:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 6749 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 84:

CCTGATGGGT GGTATGCGAG GATAACAGTTC TGAAAATCGC CGTTACTTAA TTAATGGACG	60
CGAACGTACA CCTGAGGAAT TTGCTCACTA TCGTGCGACT GGTCAATTAC CAGGAAATGC	120
AGAAACTGAT GTGCAAATGC CACAACAGGC ATCAGGTATG AAACAAGGCG GTGTCCTTGC	180
AAAACCTAGGT CGAAACTTAA CAGCAGAAGC GCGTGAGGGC AAGTTGGATC CTGTTATCGG	240
ACGAAACAAG GAAATCAAG AACACATCTGA AATCCTCTCA CGCCGCACCA AGAACAAATCC	300
TGTTTGGTC GGAGATGCAG GTGTTGGTAA GACAGCAGTT GTCGAAGGTC TAGCGCAAGC	360
CATTGTCAAC GGAGATGTTG CTGCTGCTAT CAAGAACAAAG GAAATTATTT CTATTGATAT	420
CTCAGGTCTT GAGGCTGGTA CTCAAATACCG TGGTAGCTTT GAAGAAAATG TCCAAAACCTT	480
AGTCAATGAA GTGAAAGAAG CAGGAAATAT TATCCTCTTC TTTGATGAAA TTCACCAAAT	540
TCTTGGTGCCT GGTAGCACTG GTGGAGACAG TGGTTCTAAA GGACTTGCAG ATATTCTCAA	600
GCCAGCTCTC TCTCGTGGAG AATTGACAGT GATTGGGCA ACAACTCAAG ACAGAACACCG	660
TAACACCACATC TTGAAGAATG CTGCTCTTGC TCGTCGTTTC AACGAAGTGA AGGTCAATGC	720
TCCTTCGGCA GAGAATACCT TTAAAATTCT TCAAGGAATT CGTGACCTCT ATCAACAAACA	780
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CATTCCCAA CGTAGCTTGC CAGATAAGGC TATTGACCTT GTGATGTAA CGGCTGCTCA	900
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AATGGAAGCT TCAGATATCG AACGTTTGAA AGATATGGCT CATCGCTTGC AAGACAAGGT	1200
GATTGGTCAA GATAAGGCCG TAGAAGTTGT AGCTCGTGC ATCCGTCGTA ACCGTGCTGG	1260

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TTTTGATGAA GGAAATCGCC CAATCGCAA CTTCTCTTT GTAGGGTCTA CTGGGGTTGG	1320
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AGCAGGCTAT GTGGGTTATG ATGACAATAG CAATACCTTA ACAGAACGTG TTCGTCGCAA	1500
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AGAAGATGCC GATAAACAG AATTGATGGA CCGTTTGAA CCCTCTTCC GTCCAGAATT	1740
CCTCAACCGC TTTAATGCAG TCATCGAGTT CTCACACTTG ACTAAGGAAG ACCTTTCTAA	1800
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TCGATGGTTG TCCCTGGAC ATCTTCCCAA GTAGTAGTTT CAGCGATTTC CTTGAGCGAA	2520
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TAAGCGGTG TATTGGTAGT ATGGGTCAA GGTTACGTTG ATACCCAGTT TACGAAGGAC	3000
ATTCTTGCTC TCATCAGTCA AGATGATGGT TGAGTGGCT TCGCTTCCCT TGAGGTTGCC	3060

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GAGTTCTTCC ATAGCGCGG CAGCATCAGG ATTTCTGTA GCTGTGATAG CAA GTGCAAT	3120
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GCAAGAAACA	GCCGGATCTC	AACTGGAAA	ATGAAAAACT	TCGCCAGAAA	ATTTATGAGA	5100
TGATGAACCT	CTGGATTGAT	AAAGGTATTG	GTGGTTCCG	TATGGATGTT	ATTGACATGA	5160
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AGGAATGAA	TCAGGGCACC	TTTGGAGATA	AGGATCTCTT	GACAGTAGGG	GAGACTTGGG	5280
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CAATCTGGGG	AAATGACCAA	GAATACCGCG	AAAATCTGC	CAAAGCCTTT	GCAATCTTAC	5580
TTCATCTCAT	GAGAGGAACT	CCTTATATCT	ACCAAGGTGA	GGAGATTGGG	ATGACCAACT	5640
ATCCGTTGA	AAACACTGGAT	CAAGTAGAAG	ATATTGAATC	TCTCAACTAT	GCGCGTGAGG	5700
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ACAATGCCCG	TACCCCTATG	CAATGGGACG	AGAGCAAAAA	CGCTGGTTTC	TCAACACCTC	5820
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AGGCCCTGACT	TTTGCATCTG	CTTGCTTTT	ACCCAAATCA	TTCATACCTC	TCTCAACTAG	6420
ATGTAACCTA	CAAAACCCCT	GACCTCATGA	GCCACTTTCT	TCCTCCTCAT	GAGGTCAGTT	6480
TTACTTTCTG	CTGTTCCAGT	ATCGTTTTTC	CTCGCTAGAT	TTCCCTAAAA	GGGCAGACTC	6540
CTCCCTTGGT	GCGTCACACG	ATTTTTTCAT	CTCGACTGTT	CTTTAATGCA	TCATTAACGA	6600

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CGCTTTCTT CTAGGTGGTT CATAAGGAAC AGGAAGATTG AGGTTGACTT TTCTAATCCT	6660
AGAATAAAAGT GCTGAAAACA ATTCGGAATA GCCATAGAGA CTAGACAATT TGAGGAGCTG	6720
CTTGCCTCCT GTTCGAACAC ATTTTCCGG	6749

(2) INFORMATION FOR SEQ ID NO: 85:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 1842 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 85:

TCTACCCATG GACTTTGAGG CATTCAATTGT TCCATCTTCT AGTGGCGAAT CTTTGATAC	60
AAACGATTCA ATTCACTTGG ATAGTGAAC TCTCCCGCAA ACATTTTCT GGTTAACTCA	120
ATCCAGCTGA TATTCTTTC AGCCAAAATA ATGGACAAGT TCTCCCAAAA TCCTTCAGCC	180
ATATTGCTTC TCCTTTAGTT AGATAAAATAA TGTGTTGCG CCATGTAAAT CAATTGTTTC	240
GTATCTCTG GCAATAGAGC TCTAGCCTCT TCCAAATTCA GACTTGGATA AACTCGCTTA	300
TTTGAAACCG CAAGAGGAAG TCTGATGGTT AGTTCAGGAT TTTTTAAAAT TATCTCAACG	360
AAATCCGTTA ATCTTAGATT GTCACGGTTC TAAATCTA ATAAATTGGG AGATAAAAAC	420
TCAAAACAAAT CTGAAGAATA GCTCATCATC TCAATTAAATT TGTCCCTTGT CATTTCAGAA	480
ACTGAATGAC AAGATACCTC TATGCCATAG TTTGGAAGA AATCTAAAAG AAGTTGATTT	540
CTTTGTCTAT TTTTACTTAG ATAGAGATCA ATCATGGGAG ACCTCCAAA GATTCCGGTTC	600
CATTTGATAT TCTGACACGA TTAAGGAATC TAATAAATTAGGAATCTAA TAAATTTGGC	660
AAGTTAACG GTTCTTGTGTC TTCATCATAA GCTTTTACAG TTACTTGGGT TGTAAAGTATT	720
CCCTCTTTTC CCTCGGCTCG ATAGCCTGT CCATATAAAA CAAAAACGAG ATTTTGATGA	780
TCATCTACAA AGGCATCAAC CCCATTCTT ATGTCTTGAC TTTCAAGGA TTCCATAACG	840
TTTTGAAGAT AGGATTCGTA AAATAGTGGG TAGTTATGTT TTTTATGGTA ATCATCTAA	900
AATGTCACCTT CAAACTCACA TGGAGAGTAA TTTTGACTTT GAACAGCCTA AAAGTGCCAT	960
CAAATTGAA TTGGAATAAA TCAAATAAT AGCCCCATCC TCATCAATCC AACCTTTGCT	1020
CAAAGACAAC TCCAACCGAT CTTTTAAAC TGAGTAAACC ACCTTAACCT CCAGTTTCAT	1080
ATTCTTATAC CGTTCACTCT CAAATAAAAG TTTGGGGAGC TTATAATAAC GCTCTGATGT	1140
CTGATATTGA TTAGCGGTAAC TACGCTTCAT TATTGTCCTT CCAAGACTAA AATTCCAACA	1200

	682
TTTCCAAATT CATCAAATCG GATTAACCT ACTTGTTCCA TTTCATCAAC TAACTGAGTT	1260
GCTTTACCC AAATCATTCA TACCTCTCTC AACTAGATGT AACTTACAAA ACCCCTGACC	1320
TCATGAGCCA CTTCTTCTC CCTCATGAGG TCAGTTTAC TTTCTGCTGT TCCAGTATCG	1380
TTTTCTCTG CTAGATTCC TCAAAAGGGC AGACTCCTCC CTTGGTGCCTG CACACGATT	1440
TTTCATCTCG ACTGTTCTT AATGCATCAT TAACGACGCT TTTCTTCTAG GTGGTTCAT	1500
AGGAACAGGA AGATTCAAGGT TGACTTTCT AATCCTAGAA TAAAGTGCTG AAAACAATT	1560
GGAATAGGCA TAGAGACTAG ACAATTGAG GAGCTGCTTG CGTCCTGTTG GAACACATT	1620
TCCCCACCACG TGAAGAAAAA GATGGCGGAA CGCTTGATT GTTAAAGTTT GGAAGTCACC	1680
TCCAGCTAGA TGTTTGAGAA AAAGATAGAG ATTGTAGGCG ATACAGCTCA TCATCATACG	1740
AACTTCGTT TTGATTAAGG TTGAACATATC CGTTTATCG CCAAAAATC CCTCCTTCAT	1800
CTCCTTGATG AAATTCTCGG CTTGACCAACG TCCACGATAA AG	1842

(2) INFORMATION FOR SEQ ID NO: 86:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 19390 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 86:

TCATCTTAT CTCCTCGAAA TTTCTAAATA TAGCCATTAT AACAGAATT TGTGAAAATT	60
CCTATTATAG TAAATCACTA TTTCAGTATA AAAAGAAAAA ACGAACAGA CGATTGCTC	120
TTCTTAAAT CTGAAAATAG CTTTCCAGAA AGGATTAGCC GATTTTTGC AGATTGAGCA	180
CTGCATCGTG ACTCATCAAG ACTTGACCAT ACTCTTGTAA GACTGAGCGA CTGATATCAC	240
TATCGCTGC AAACTCGCGC ATACGGGCCA ACAGCCAAGC TGGATATGGG CTTGGATGAT	300
TTTCAATATC CACTAAAATG GTCAAATAAT AGCGCTCGTT CATTGTAG AGTCAGAAG	360
TTTCCATTTC AAAAGTCACT GTCTTGGCAA AAGCTACCAA GTCAGCCAAC TTAGCAAAAG	420
AAAGGATGTA GTAGATGTA GGTTCTTCT TACTCTCAGC TTCTTGTCA GCCTGCTCTT	480
GCTCTTCTTC CTTGACTTCA ACTTGCTCAA GAGATTGAAT GGCTTCGATA TCATCCTGG	540
TTTTGCTGC GATGTTTTT TCCAGGGTTT TGATAAATTC ATCTGGAGAC ATTTGAGCCA	600
ATTCTTCCAT ATCTGGCAA TCCGATAAGT CTTCAAATC TAGTTTTGG TCAATCTTG	660
ACTTGTCAC AAAGACATCT ACCTTATCAG CTTTGAGT CACACGGAAG CTCAACATGC	720
CTGTATCCAG AAAGCTATCA GGCATCTCTA GCTCATCCAA GATAGCATAA AAGAACTCTT	780

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CTGTTTTTC TTGAGGAACG AGAAAAGTCAG CAATCTCCAT TCCACGATCC ATCAAATCCT	840
CTAAAGATAT CGTGATTTTT AAAGTTGTAT CACTAATTG TTTCATTTTC ATTGCTAGTA	900
ACCTCATACT TTCAGTTCTA TCTATTATAC TAGATTTTA CGATTTTATC AAAAGAAGGC	960
TCCTCTATAC GGATAGATTT TCCCTAGGGT CTTTCTATAG GAGACTCCAA AAGAAAATT	1020
CTGCAGACAG ATAGAAAAAG CCTTCAAAAT CGGCTAACAG CCGACTTTGA AGACCTTATA	1080
CATCAGAATA CTTATAATTAAAGGTTGCT ACACCGAGGA TAGAACGATT TAAGTTCTG	1140
AGAATTGAA GACTTGCTC AAATTCTTA TAACGAGTCA CTCCGTACTC TTCAACAAGA	1200
AGGACTGTAT CTCTTCCAA AAGAGATGAT ACATCCTGTA AATCTACAAA ATGCATTCC	1260
TTTAAAGCTT CTTGACTCTG TTTCAATTAA TCTAAGATAG CTTTATTGTA GCTAACGATG	1320
GTCAATTCCCT GTCCAGTATT TTTGTATGAC AAAACATCTG CTAGGTTAGC AATTGTTGTA	1380
ATCTCTGTTA CAAAATCAAT TTGATACTGA GAAAATCAC CTACTCTATT GATTGTTGGA	1440
TTAAAGAGAT AACTAACAC ATTTCCCAC ACAACCAAAA TCACACAAAC CACTCCAATA	1500
ACAACCTAAAC GAAGAACATCAG ATTTTCACA TTTAAGCCAA GCGCTGTTTC ACCATTGCG	1560
TTCAATTCTT TAGAGTTGAT GGTTCAGT TTTCAATTTC TCACATTGTC ATAGGCATGT	1620
TTAAATTCT CAATCAACCC ATCAATTTC TTCTCTAACAA AGTTATTGGC ATCTTACTT	1680
GATGTAAAAA TTTTCACACC AACCCCTGCA TCGTCAATCA TATAGTAGAC GGTCAATT	1740
TTCCACCAAT AGTCATTGCT TGAATTTC AAGGTTGTTT CTGTCGTGTC TAATTCACTG	1800
GCAATTTC TCAACTCACT GGTTCTACA TCATTGAAAA GATAAGCTCC ATTCAAATTAA	1860
CCATCAATCA ATTTCCCATA AAAATCACTA TAACCACCAA TTTGATGATT CAAAATCGTT	1920
TTGTCCGACT CTTTTGGAGG AGTGATTTA TAGATAAGAT AAGTTGAATA ACTTGTGTA	1980
TCTTTGACAG TGTTTTTATT CCTAACTGCT TTAATTGTAATGGTACAGC AATGAGAGCA	2040
AATAAAGCGA TGAGAGCTAA AATATTGCT TTTCGCTTT TATAAAGATT TGCAAAACAA	2100
TCAGCTACTG AATAATGTTCA AAACATGATT TTTTCTCCT TTGTTTAGTA GATACTAGTT	2160
TTCCCTTGTA ACCATTTTG CTACAAATAT AATCACAAGA ACAATTCCCC AGAATTGCA	2220
TGTAAATAAA TTGAAGAAC TTTCTGAAAA GCTGCTTCTT GGCATAAAAGA ATAGATTATT	2280
CAAGATGAGT AGGGATAAAG CAAATAGGAT TGTCTTGAG CGATAGGCTA CTTGCAGCAT	2340
GGCTATAAAAT AATACGCCGA GTAAGAAACT AAGCAGAAAG ACTCCAATCA TACCATAGTC	2400
GGTATACAAC TCCATGATAT AACTACTTCC GATACCATGC CTTTCAAGT ATTCCATTGTT	2460
CAAGACAAGA TAGGATAGAT TGTGGGCATA ACTATTACTA TCAATAGCTA GTTCCACACT	2520

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ATTGGTTGTA TGTTCAAAGG CTTTCCCTCC GAAAATGGCT CCCAAACTCC CCCTTGCAAA	2580
ATAATCAAGA ACAGGACCAA AAGTAAAATT ACGGAAATCT CGGTAGGGGA GGCTACTGTT	2640
AAATAGAAAA CCTCGAGCCA GAACACCAAA ACTAGTCCT TGTTATAGA TAAAGTCAG	2700
TAAGATATCC CAGAAACCTG TATGGGAAAC TTGGACATTA TCCCCTACAT AATTGAGTAC	2760
TCCCCTCGCT AACATGAGAA TAGGAGAACC TACAAAATC GCTAACTTTT CTTTAAACCC	2820
AATCCATTTT CCTTTTCAG TTGCTCCCG CATAAAGTAA TAAACAAAAG CAAATmAAT	2880
ACTTAAAATA AAGGGATTC TGTCCTCAAT TGCCAAATGA ATAGTATTAG CTGCAATAAA	2940
GGAGACAAAGC ACTGCTGTGG CCTGCAATTT CTTTGGCTTG GTTGCCAGAT ACATACACAT	3000
TGCATAGACC GTAAAGGTAG ACAAAATGTA GGAAAATAA GGCAAGTTAC TTTCAAAATT	3060
TGCATAGTAG GCATAGTAGG AAGTCTGCAA ACGATACAAG AGCCGTTCAA ATAACCGAAT	3120
GAAATAGAAA GGATAAGTTA GAAGAAAAAC TCCTAGTGAT ACAAAAGCGTA ACCGCTTGAT	3180
ATAAAACCTCT TTTAGAGAAT TTCCCTATATT TGCTACTTTT ATTTTCTTCC TAGCTATGAA	3240
GTAACCGAGCC AGAATGCCTC CTGTGGTCAA GCCCAGAAC GAAATCATGA CAACTATAAA	3300
GGCAAAACGA TAGGCTATTG GATGATAGGT ATCCAAAGCA CCATCCCTAA AATAATCAAT	3360
GGTCGGTCTT GATACCAGAA ATACAAAAT GGTTAAATAG AAAATAAAAT GGATTAAGTA	3420
ATACTTGATA TCATTCAAC AACCAATTAA GCTACTAACCC AACAGAACAA ATAAAGTAGA	3480
AAAGTAAGCTA ACATTATTAT TATTAACAG ATACACAATT CCACCTACTA GCGTCAAGGC	3540
ATAACTGACT ATGGTCAAC TAAATAATAA TCGTTCCCA TCAATCCTT GGTCACCCCC	3600
GTTCTAATGT AATTTTTAG ATTTTCAAT ATTTTCAGT AATAAGAAC GATATAAGGA	3660
AATATTTATG AATAGGGCCA AACGACTAAT TCTTCTCCCC TTACGGAAAA TTGGATTCCCT	3720
AGAAAATAGCA AAGGCATGGC CTTTAAAAAA ACGATGAATC TGAGAACAGG CTTCAAACGT	3780
TTTATACTGA TCATCTAGCA ACATCTTATC CAGAATAAG AAGTGGGCAT AGGCAATCT	3840
GAAAAAAAGCG ACCTCTTCAGATA AGTCAGGATA GTTTTCACA ACTTCATTAT AAAACTTTG	3900
GTAGATATCA ATATAGGCTA AATCCTCTC TGCTAGGGT TTGGTCGTA TACTATCCCC	3960
TCTATGGAAA TAGTAATAAT AGGGTTAGT ATTAACCACA TACTTCTTGG CCAACTTGAT	4020
TAAATCAAAA TGGTAATAGG CATCTCGTA AATCAACCCCC TTAGGAAAGG ATAGGGCAGT	4080
TGCAATCTGT CTCTGATTA GCTTATTGCA AATCGTCCCA GGTATTTTTT CACCTATGAG	4140
GTATTCTTT AGAAATGTTT GAGAACATCAC GACAAAATAG TCATCCTGAT TGGCTGACTG	4200
TGGGCTTCAGA TCATTAGCAT AGACATTCAAT GACACCACAG CTCGAAACAT CCGCATCTTC	4260
TTGAACTAAT TGCTCATATA AGCTCTGAAT CATTCTGGAA TGGATATAAT CATCTGAGTC	4320

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AATAAAAATC AGATAATCCC CGTGAGCCTG CTTCATCCCA TCATTTCGTG CTTGCGACAA	4380
TCCTTCGTTT TTTTATGAA GCACTGACAC CCTGTCATCT TGTTCAGCGA TTGAATCACA	4440
CAAGCGACCA CTTTCATCTG TTGACCACATC ATCAACAAGA ATAATTCCA GATTTTGATA	4500
GGTCTGCTTC TGAATGGAAG CTATCGATTT TTCTAGGTAC TGCGCCACAT TATAGACTGG	4560
CACAATCACA CTAATTAATG CAGTTCCAT GCTACTCCTC TAATAGTTTT TCTACTTGGT	4620
CGATTGTGTT TGTAATTGTA AATTGTTGAA TGAATTGGCT AGCCTCATCG ACATCAAAGT	4680
TTGAGGCAGA AGTCATGTA TTAGTAATCG CCTGAGCTGC CTCTTGATTC CTCTCAATGA	4740
TTTGTCCAAA TCGTCCTCT TGGGATAATT CCTCAGCCCC TCCAAACGTCC GTAGAGATAA	4800
AAGGGAGTCC CAGACTCAAG GCCTCCACAT ACACCTCCAGG AAAACCTTCT TGTTTAGACA	4860
TAGACAAAAG AACTTCGTC TGAGATAGAT ACTGATAAGG ATTTTTTGTA TAACCAAGGA	4920
AATGTACATA GTCCTCAATC CCATACTCTT TGACTCGTTT TTTCAGTTC TCTTCCATAT	4980
CACCAAGCCCC GATAAAATAG AGATGATAGT TTTTCCCTC TTGGTGTAAAT AATCGTATCA	5040
CTTCCACTAC ACGGTCAAGA CCCTTATTTT CCTCAATCCG TCCGATAGTA CAGATACTTT	5100
GAGGAGCAAT CTCGATATCG ATCTTCTCTT GAGATTTTC TAGAATAGTC TGAAAATCAT	5160
ATCCATTGTA GATTGTCTGT AATTAGAAG TATAATCTGG ATAAACCTTC TTGATAGAAT	5220
TGCTGGCTT TTTGAAATC CCTACAAATTG TATTCCGAGC ATCCAACCTGG CTTCATGTG	5280
ATTCTCTTTT AGAGCTATCC TTAAGAAGTT CTTCAAACT TCCATGAATC CAAGATATCT	5340
TCTTGACTTC TCTCTTTTA GAGAACACA GTGGTGGATT CATAATGGTA AAAGAAACTT	5400
CAACATCATA ATCATCTTT ACAAGCAAC GACGAGTCAG TCTTGGAAAA TAAATTCTCA	5460
TTCTCCACAA AAAAGCTCGT AACCATCTGG TTTGGCGATA ATCTTGAAGG GATTTAAAA	5520
TGCGTACATG CTTTGGAAACA GATTCAATTC CCTTGTCAA GTGCTCCATT TCAAGAATAT	5580
CAATATCATA CTTTCTGGA TCCAGATTTG AAACAATGGT TGATAGAATC TTCTCTGCAC	5640
CACCTCCAAG AGAAAAGAC CACATAAAAA ATAAGATTTT TTTCTTAGCC ACCATATTCT	5700
CCCTTGTATT CTGTATAAGA CTTATCCATA TCAGCGATGA CAGCATCATG ATGGGGTACC	5760
TGCTTGCTG CTGGTGGAGG CGTCATATAA TCCCCAAAG CAGTTCTGAG ATAGACATCA	5820
TAGCCGATTG GAATAGGCAT CTCTGTTCC TCAAATGGCA AGAAAAGATT GTCTCAAA	5880
GATGTGATTG GGTACTTGTT TCTCATGTAG CCAGGACCTG AGCATAATTC TGTAATGCCA	5940
TCACAATCAG CCAAATCATA CTTAGTCATT TCTTCTCAG CTTTTTCCA GATGCGATAA	6000
CGGAGAGATT TTGGAGTCAA ACCCAGTAAA ATGCGACTTC CCCATTTCAT GAGATCACCA	6060

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TGCTTTCTG	GAATAGTTG	CGCACAAAAG	AGTGAATAAA	TCAAGGCCA	ACGAACCTGT	6120
TTTTCCGCT	CAGCTGGATT	TTTCGGATAA	TAATCCAAG	GCAAAACATC	CAAGGCCAGA	6180
CCATGTGGCA	AATCCAATC	CTGCTGATAA	GGCTTGATAC	AGGTGGTTT	CTTGTACCGA	6240
ATGGTAATAA	AAAGATTACG	ATCAACAAAA	TCCTTGAC	TCTTGACAA	GAAATAACGT	6300
TCATCTGCAT	AACGAGGCCA	TAATTCTGCT	AATTTCTCAT	AATCTTACG	AGGCATAAAA	6360
AAGTCTAGGT	CGTCGCTCCA	AGGAATAAAAT	CCCTTGTTTC	GAAGGGCACC	AATAGCGCCT	6420
CCGCCACAGA	GATAACAGAG	CAAATCATGT	TCTTTACAAA	AGGCCACAAA	ATATTCAAGCC	6480
ATCTCCAGAC	TACGAGCCTG	AATTGCTTT	AAATCAGTC	TATTGTTCAT	TATTCTTCT	6540
ATCGTATCGT	TTCATTATAC	CACAAACAAG	GGGTGAAAAT	CTATTGCAGA	CTGTAAAAAA	6600
TCAAAGCCTG	ACTGCTATCC	AAATAGCTAT	CAAACATTGA	TTTTCTGTC	TTATACTCTT	6660
CGAAAATCTC	TTCAAACAC	GTCAGCTTCA	CCTTGCCGTA	GGTATAGGTA	ACTGACTTCG	6720
TCAGTCTTAT	CTACAACCTC	AAAACTGTGT	TTTTAGCAGC	CTGCGGCTAG	CTTCCTAGTT	6780
TGCACTTTGA	TTTCATTGA	GTATTATCTT	ATCTTAAGCC	CATTGAGCG	AGCTTGGTTT	6840
GATATTGTT	TTGATCAACC	AGCAGGCCA	AGCCCCATA	AACATCATAG	GCATCTACCC	6900
AGTCACCCAG	TTCTGGAATC	GTCAATTTTT	CAATACCATT	TTTGCTCCA	TCCAAAACAG	6960
ATAAACCGTT	TGTTAGGAGG	AAAGTATAGG	GTACGTTGGT	TGAGGTCTA	GCAAAACCT	7020
TTCCAAGAGC	TTCAGAACCA	GTGAAAAGTT	TAGTGGATC	TTAATTGTC	TCTAAAATTG	7080
CTGTTAAAC	TTGTTGCTGT	CTTTTTGTAC	GGCCGTAATC	TGCCTCATCA	TCATCACCGA	7140
AAAGACGATA	ATTGAGCAGG	GTCGAGCCAT	TCATCTGCTG	TTTCCGACT	TTAATGGTTT	7200
GGGTTGGAGA	CTCAGTCTCG	GTAGCGTATA	AATCATCTCC	GACTGTAGCT	TCTGTTAGGG	7260
GACGCCATT	CAATGTTGAA	AATTGAGCAT	CAATCGTCAC	CCCATCAGGG	AAAAGCGTGT	7320
CAATCGCTGT	GGCAAAGGCC	TGAAATCAA	CCAAGGCGTA	GTACTTAATG	TCCAAGTCAA	7380
AATTATCTTT	CAAGACTTGG	CGAACCATTT	CTGCCCTTT	TTGCCCTCT	TGTTCTCCTA	7440
ACTCGTAGGC	TACGTTAAC	TTGTTATCTG	TCTGTTTCT	ACCATTAATC	ACTTGACTAT	7500
AACCACATAT	ATAGACAAA	TTATCACGCA	TGAAACTGAC	TAGTTTCATT	TTCTTATCTG	7560
AGCCCCCGAC	ATTTAATACC	ATAATAGAGT	CAGTTCGTGT	CTCAACACTG	TTCTGGCCGA	7620
TTCGACCATC	AGTACCCATG	ATTAATATAT	TAACTCCATC	TCTAGTGTCC	TGACCATTA	7680
AGACTTCTAC	TTGAGCTGCC	CGGGCATCAG	CAGTTTTCTT	TGCGCTAGCA	TCTTGGTAAC	7740
CACGAAAAAA	CATGAATACC	ATGGCCAAAG	CCACACAGAC	CAAAAGTGAA	AAAATCACCA	7800
TAAAAAATCG	TTTAAGACGG	AGCTTCCGTC	TTTTCTTTT	TGGAGGGAAA	GAGAGTGCTT	7860

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GTGATTGGAA TTGTGAGCGA CTCCGGTTCG CATAGCTTGG TAAGTCAACC TGCTCTTCTC	7920
TTTCTTGTTC CAAGCTAGAG CTACTATTC CCCTAGCAAG AGTTAGCTTT TCTTGCAAAT	7980
AGGCAAACTC ATTTTTTCT CTCTCATTGA GATAGTGAAT ATTTTTAGC AAATAATCAT	8040
AACGCAACTG CTCATGATGA CTTAAGGGAT TTCTTTACT CATCTTCTCT CCTTTCCATG	8100
GTCTGATATT GGATAAAATAG GATAGGCACC CAGAATTATA TACTGGATTC CAATCGCTTC	8160
TAATTCTTTT TGGGCAAAGT GGACCAAGTC CTTATCGGTA TAATCCACAT CGATAATGAA	8220
AAAGTATTCA CCCAGTGCTG TCTTGACTGG ACAACTTCA ATTTTGTCAGTCAATTCC	8280
TCGCCAAGCA AAGGTCGACA GGGCCTTATA AAGTGCACCT GGAAGGTTGT CAGGTAATGTT	8340
CAAGGCCAAA CTCATCTTTT CAGTTTGTC TTGCAAGGGAA ATACTAGGCT TTTCAGCTCC	8400
TAGAACCCAG AAACGTGTGA AATTGGCTTC CATTTCCTGA ATATCCTCGG CAATCAGTT	8460
CAATCCATAT TCTTCAGCAG AACTTCTAGG TGCAACTGCT GCAAAGGGCT GGTCTGGATG	8520
TTCGAAATA AAACGGGCCG CATAAGCTGT ACTAGCTGTT ACCTCGATTT GAGCCTCTGG	8580
ATATTGTTCA TCGATGAATT TCTTCTTG AGCCAAGGCC TGTGGATGTG AAAAAATCTT	8640
TTCAATCTTA GTATGGCCTG GAACCACCAT CAACTGCTGA TGAATAGGCT GAACGATTTC	8700
TGCTACTGCT TGGAATGTGAG CCTGATGAAA AAGATAGTCC AAGGTTTCAT GAACACTACC	8760
CTCAATAGAA TTTCAACTG GCACCCACAGA ATAGTTCACT AATCCTTGCT CATAAGCCTT	8820
GATGACATCT GTAATGTTGG CAAAAGCCTG CAATTCCCTCA TGAGGAAAAG CTGTCCTGCAC	8880
AACGTGGTGT GAAAATGATC CCTTGGGACC TAGATAAGCA ATTTTCATCT TAGTTCCCTC	8940
ATAATTCCCT CTGGGCTTAG CTTGGTCACA TCCAAAACCC GACTAGCCAC TTCCCTCATAC	9000
CAAGCCTGTC TTTCTTGGAA AATAGCTACT AGTTCTTCCT TGCTATTATT TAGAAAAAGC	9060
GGTCGCTGAT TGTCCCTTATC AGCTGGCATA CGTTGGTAGA GGGTTTCAAA ATCTGCTCTC	9120
AGGTAGATGT TATCTGTATT AGTCTTGAGT AAGTCACGAT TTCTCTGAGA AATAACCACT	9180
CCTCCTCCAG TTGACACGAC TTGGTCTGTT TGTAGTAAAT CAGCTAGGAC TTCTGATTCT	9240
ACCTGACGAA AGGCTGTTTC TCCCTTTCA CGGAAAAAAT TCGCAATGGA CATAACCTAGG	9300
CGATTCTCAA TCAGAGCATC CATATCAAGG TAATTAGGGT CCAAGCCTCT TGCAATAGTC	9360
GATTTCCAG CCCCCATAAA CCCTAATAAC ACCTTAGCCA TGAATCAAGC TCTCCAAATC	9420
ATCAAAGAAA CTAGGATAGC TGGTATTGAT GGCTTCTGCA CGGTCAAGCT CCACCTCTCC	9480
ATCTGCAACC AAGAGGGCTG CGATAGCTGT CATCATGCCG ATACGGTGGT CACCAAAACGT	9540
ATTGACTCTA GCACCGTGAA GAGCTGATTT TCCTTGATA ATCATCCCCT CTGCCGTAGG	9600

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AGTAATATCT GCTCCCATA C TATTAAAGGC GTCTGCCACA ACCTGAATAC GGTCTGTTTC	9660
CTTGACCTTG AGCTCCTCA G CATCCTTGAT AACTGTTACA CCTTGGGCTT GGGTCGCAAG	9720
CAGGGCAATA ATGGCAATT CATCAATCAA TCCTGGAATC AAAGGCCAC CAATCTCTGT	9780
TCCCTTCAGA TCAGAAGACT CAACAATCAA GGTAGCAGAT TTAGCGACTG GATCGATTTC	9840
AGTTATTTCC AATTTCAC CCATGGCACG AATGACATCA ATAATACCGG TGCGAGTTTC	9900
GTTGATCCCC ACATTCTGCA GCACCTAGACG AGAATTTGGA GCAATCAAAC CTGCGACTAA	9960
CCAAAAGGCT GCACCTGGAA TATCTCCTGG TAGGACCAAC TTCTGTCCTG TCAATTTTG	10020
TGGCCCTGG ACTGTGATT TCTTACCATC CACACTTAA TGACCACAA ATTGTTCAA	10080
CATATCTCA GTATGATTAC GGGTGTACTC TTTTCGATA ATAACGTACT CCCCTTAC	10140
TTGTAAGGCT GCAAACATCA AGGCTGACTT GACTTGGCA GAGGCAATTG GCAACTCATA	10200
ATGAATAGGT CTTAGGTTTT TCGTCCCTTT TAAGCGAAGG GGAGGCAAGT CTCGTTCACT	10260
TTGCCCTGAA ATGCTGACGC CCATTTTTT CAGTGGAAAGG GTCACACGGT CCATAGGACG	10320
TTTGGAAAGA CTATCATCTC CAAACATCTC TACTTCGAAA TCTGCACCAAG CAAGGACACC	10380
TGAAATCAGG CGAACATCGAGG TGCCAGAATT TCCCATATTA AGGGCATTGTT GTGGCGCTTT	10440
TAAGCCAGCC ATGCCTACAC CTTGAATGGT AAAAACCCCA TCTTTATCCT CAATTTCAAC	10500
ACCAAGGTCA CGAAAAACCT GCATGGTCGA AAAAACGTCT TCACCTCGCA GAATATCATA	10560
AACCTGGTC TCACCCCTAG CCAAACCTCC AAAGATAATG GAACGGTGCG TGATAGACTT	10620
GTCACCTGGG ACGCGGATAC TACCATGTA ATGGCGAATG TTTGTTTTTA GTTTCATACT	10680
GGACCTCATA CTTGCAATAC TTTTACCTAT TTTATCATAA AAAGCCAGAA ATTCCCTTAA	10740
AATTCTGAC TTTAGGATCG TTCTTTCTT ATTCAGCAA TTCTGAAACT GGTTCAAAAA	10800
CAATTTTTC AATATCAGAA AGGTAAATGG CCAATTGTTG TTGCTTGGTA AAGAATTCTG	10860
ACAAGGGCT ATTCCTTGA ATCTGTTAC CAAAGCCTTC CATCTTAGCT TGGAAGGACG	10920
CATCTGGCAT TTGACCTGTC TGTGCTAGTT TTTGAATTTC CTCTTGAAAG GCAAGATAAT	10980
CTGTAAAGAT TTTGCTTGCC TCAGCATCTG CTGCAATCGC ATCTTTAGCT GCTTTAACAG	11040
CCTTGTATTC TGGTAATCCG CGTAGACCGC GACTGAGTTC GTTTGCACCA TCGTAAATAT	11100
TTGACATGTT CTTCTCCTTA TTGATGACG ACTGTATAGT CAGTATTTTC TGTTATGAGA	11160
TGCTCAGCTC TTTCCAAGTC TTGAGCATTT TAAATGAAA TTGAGTGGAT TCCGTGAATA	11220
TCCTCACGAT TTCTCGTT GATGTGGATA TAAACCAAGG AAGTTCCACG TAGCAGTTC	11280
AAAATCCGCA GGATGACATC TTCTTCATCA GGAACGTCAA CATAGAGGTC GTAAGAGCTA	11340
TCCACACCCAC CACGCTTATG GATTTCCATG GTCTGGCGTT GTTCACGCGC TTGGTTAAA	11400

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AAGTTCCAAA	TTTGCTCTTC	ATCTCCCTTA	CTAATGGCCT	GACCAATCGC	TTCCAAACGT	11460
TCCCTGAAAT	CCTCAATTCT	ATCCAGAACG	ATCTCGCTAT	TGGACAAGAG	AATGGAGGTC	11520
CACATTCTG	GCTCGCTTTC	CGCAATTCCGG	GTCATATCTC	GAAAACCACC	TGCCGCAAAG	11580
CGCCTGCCA	TCTCATGCTC	TTGAGCATAG	ACCGCAGTCT	GCTCCATGAG	ACTAGAAGCC	11640
AAAATATGAG	GAAAATGGCT	AATCTGAGAA	GTGACACGAT	CATGCTCCTT	GGCATCAATC	11700
TCGATAAAAC	GAGCATGAAG	ACCTGAAAGC	AGATCCTCA	TTTCCTTAAG	CGTGTCTGGA	11760
CTTGTCAAGGC	TTGAAGGTGT	AAAGATATAA	TAGGCATTTT	AAAAAGATT	GACATCTGCC	11820
GAAGCAGCCC	CTGTCTTGTG	ACTACCACCC	ATGGGATGGG	CCCCGACAAA	GCGAACAGAC	11880
TTGCCAGCCA	AATACTGCTC	CCCCGCATCC	ACAATGGTTG	ACTTGGTCGA	ACCAGCATCT	11940
GAAATAATAA	CGCCTTCTCG	CAAATCCAAA	TTGGCCAAC	CCTTAATGAA	AGCAATAGTT	12000
TGTTTGATTG	GCAAGCTGAG	GATAATGACA	TCTGCCAAAG	GAGAAAAC	AGCAAAATCA	12060
TCCGTTGCAC	GGTCAATCAT	ACCTTCTTC	AAGGCGATAT	CTCTCGAAGC	TTGACTACGA	12120
TTATAAACCTA	AAATTTCATA	ATCTGGATGA	TCGCCTTGA	TACCAAGTGC	CATAGAGGCT	12180
CCAATCAACC	CAAGACCTGC	GATATAGATT	GTTCGGCCA	TAGGAAC	TTAATAGTTC	12240
TTTGTATAGT	CTCGGTGTTT	GGCTTACCGCT	TCTTTAGTT	CCTCAAGATT	ATCTGATGAG	12300
AATTTTCGA	GGATTTCTTG	CGCCAGAAC	GTGCTACAA	CTGCTTCCAT	GACCATTCC	12360
GCAGCTGGAA	GAGCAGTCGG	ATCACTTCTC	TCCACGGTTG	CCTGTGAAGG	TTCTGGGTT	12420
TCGATATCCA	CACTCATAAG	AGTTTATAA	AGAGTAGGAA	TGGTTTCAT	GACCCACGA	12480
ACAACGATGG	GTTGCCATT	AGTCATACCA	CCTTCAAAC	CACCTAGATT	ATTGGTACGG	12540
CGAGTATAAC	CGTCTTCTTT	AGACCAGAGA	ATTCATCCA	TAACCTGGCT	GCCTTACGA	12600
TAACCACGCT	CAAAGCCAAG	ACCAAATTCC	ACCCCTTAA	AGGCATTGAT	AGAGACAACA	12660
GCTTGAGCCA	ATCTTGATC	CAATTTCTA	TCCCATTGGA	CATAGGAACC	AAGACCAACT	12720
GGAACGCCCTC	CGACGACTGT	CTCCACAACC	CCACCGATGG	TATCACCATC	ACGTTTGATT	12780
TGGTCAATAT	AGTCCTTGAT	TTCTGTTCT	CGTTCTTGGT	TGACAATAGA	AACTTCAGAC	12840
TGGGCAGCTC	TTTGCTTAAT	TTCAAGCAGT	GTCAGATTTT	CAGGAACATC	GATTTCTTG	12900
CCACCAAAGA	CCACGACATG	GTGGCAATC	TCCATATCCA	GCTCAGCCAA	GAGGCCTTGT	12960
GCTACTGCAC	CAACTGCCAC	CCGCATGGTG	GTTCACGAG	CTGATGAACG	CTCCAAAGAA	13020
TTTCGCAAAT	CATCAAAACG	GTACTTAATC	CCCCCAACCA	AATCGGCATG	ACCTGGCGA	13080
GGATGAGTAA	TTTCCGCTT	GCTTTTAAGG	CGGTCTTCAA	TGTCTCCGC	AGACATGATG	13140

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TCCAGCCATT	TCTGGTGGTC	CTTATTGATG	ACATCCATAG	TAATAGGCCG	CCCTGTGCGTC	13200
TTCCCGTGGC	GAACGCCCGA	AGTAAAGACA	ACCTGGTCAT	TCTCAATCTT	CATA CGACCA	13260
CCACGACCGT	AGCCACCCCTG	ACGGCGTCTA	AGGTCCCTCAT	TGATATCCTC	AGCTGTCAAT	13320
GGAAGTCCAG	CTGGAATTCC	CTCAATAATA	GCTGTTAGAC	GGGGGCCGTG	TGATTCTCCT	13380
GCAGTTAAAT	ATCTCATACA	CTCTCCTTAT	TTTACCAAGT	AGTCTTCAT	CTCTTCCAGA	13440
GAAACTGGGT	GAATGGTCGC	TGAACCAAGC	TCTGGCACCAC	AGACCAATTT	CAAGGTGTTA	13500
CCACCGCGTT	TCTTGTCTATG	AGTAAGAGCC	TGATAAAAGCT	TGCCAACTTC	CCAATTTCA	13560
TAGTCAACAG	GCAAACCGAA	TTTCTGACAC	ATCTCTGTGA	TAGATTGGGT	AATGCCAGCT	13620
GGCATGAGGC	CTTTTCCTC	AGCAACCTTG	GAAATCTGTA	CCATTCCCCT	GGCAACAGCC	13680
TCTCCATGCA	TGACCTTGCC	ATAACCGGCA	GTCGCTTCGA	TGGCATGGCC	AATAGTGTGG	13740
CCAAAATTGA	GGTAAAGACG	AATACCATTG	TCCAACATC	CTTCAACCAC	CATCTTGCCC	13800
TTCACCTGAC	AAGAATGTT	AATCAAGGTC	TCTGCATGTT	CCAAAATACT	CTCAACAGAA	13860
CCATTTCAGTC	CCGTCAAGAG	AGCCCACAGT	TCTGGATCCT	CAATCAAGCC	ATACTTGATA	13920
ACTTCACCCA	TCCCCTCAAT	CAACTCTCTT	TTTCCGAGGG	TTTCAAGAAC	AAGTGGATCA	13980
ATCAGAACCC	CATCTGGTTG	GGCAAAGGTC	CCCACCATAT	TTTAGCAAA	TGGGTATTA	14040
ACGCCTGTCT	TTCCACCGAT	AGAAGAATCA	ACCTGAGCTG	TCAAACACTG	CGGAATCTGA	14100
ACAAAGTGA	TACCCCGCAT	ATAGGTAGAG	GCTACAAATC	CAGCCAGGTC	CCCAACAAACG	14160
CCACCACCAA	GAGCAACGAT	TCCATCGCTA	CGAGTCAGAC	CTTGCTTGAC	TAGAAATTCA	14220
TAGACTTTCT	GAACAGTAGT	TAATTCTTT	CTTCTTCAC	CTTCTAAGAA	ATCAAAAACA	14280
GCTACCTGAA	AACCAGCATC	TTCTAGGCTG	AGCTTGACCT	TCTCTGCATA	GAGAGAGGCT	14340
ACATGGTTAT	CTGTCACAAT	GACTACCTTT	TGCGGTTGCC	AGAGTTCTCG	CAACCACTGA	14400
CCAGCCTGGG	CCATACAACC	TTTTCAATC	TGAATATCAT	AAGGATGGTG	AGGAATATCG	14460
ATTCTGATTT	TCATAGGAGA	GTCTCCCTTT	CTTTATGGT	ATTTTTCTGT	TAAAGACTGC	14520
CAAATCTCTT	CTGTCGGCAT	TTCTTGCCT	GTCCACAGTT	AAAAAGCTTC	TGCAGCTTGA	14580
TAGAGTAAACA	TTCCCAGAAC	ATTGACTGCT	GGATTGCCCT	GACTTCTAGC	CCATTTCAAA	14640
AACGGTGTTC	CAAAGGGTTG	GTATATGATA	TCTGCAACTA	AAAGAGTTTC	TGGTAAGACT	14700
ATGTTTCAG	GAACAGGAGA	GGATTGGCCA	TCCATGCCCA	CACTGCTGGC	ATTAACACTAGC	14760
AAATCCGACT	CGGCAATCCT	TGCTTGCAGT	TCAGAAACAT	ATTCTAAAGC	ACACAAATCC	14820
ACTTTAAAC	CTGTCGTCTC	CTGTAACCTG	TCTAGGTAAG	GTCTTGTGTTT	TTCCATAGAA	14880
ACGGAACGAA	CAAAGACCGA	AATCTGACTG	ACGCCATCCA	AAATAGCCTG	TGCCAAGATT	14940

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GATTTAGCCG CACCACCTGC ACCCAGCAGG GTCATCTTTT TACCTGAAAT TGTAAAAGAA	15000
GGCAAGCACT TAAAAAAATCC CTTGCCATCT GTATTATATC CAATTAAATT GCCATTCTCA	15060
TTGACAACCG TATTAACCGC ACCAATCAAG CGCGCTTCAT CGCTCAGCTT ATCCAATAA	15120
GGAATCACCT GCTCCTTATA GGGCATGGAC AGATTGATGC CAAACATCTG GTAGCGACGA	15180
ATATTGGCCA CTGTTTCTAC CAAGTCACTC GCTTCAATCT CCCAACGCCAC ATAAGCACCG	15240
TTGGTAGCTG TCGCCTCAAA GGCTCTATTG TGGATGAAGG GAGAAATAGA ATGCTTAATA	15300
GGATTGGCAA CAACTGCAGC TAAACGTGTA TAGCCATCAA GCTTCATCCA AAATCTCCCT	15360
GATTTTTTC ATGCTAGCTA GAGAAATCTG CCCAGGGCA CTAACCTCAT CCAGACTGGC	15420
AAAAGACCAA CTCGAACCGC TCACATCCGC AGTGATACGA GAGACCTTCC CCACCTTACC	15480
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CATCAAGTCT AAGACATCCT GCTCCGTGTG AGCCATCACCC GCAACCTTAA CAAGTTTTGG	15600
ATTTAGGATC GTCAACTCTG ACAAGATTC CATCATGTT TCAGGTGTTT CTTGGAAATT	15660
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ATAGTACTCA AAATCAATAT AGTCTGGTTG ATAGAGTTGC GCAACTTCCT TGATTAGATG	15780
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CAGAGCCTGA GCCTCCTCTA AACCTCTTGG CATTACTGAA ACGATTAATT TCATTTACTA	16020
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CCTCCCGAT TTAAAATCTC AAGACTCTGG GAAATCAACT TGTGATAATC CTPGGCCACA	16260
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TGCTCGCTG TGCTGAGCCC ATTTGCCTGA AAATGCGCTT GAGACAATTC TCGTGAACGT	16440
TTGGCTAGAT CAACAGAACT TGTATGGCTA GCTCCTCCCA TGGCCGCAGC TACTGAAAAA	16500
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ACTTGATACA GGACACCATT TTCTAAAACA TTGAAAAAGT CAGGTGCTTC TTGACCATAA	16680

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ACATGGGCAG ATTCA TAGTC CAAAC CCTTA AAGCGGATT TCTCATAAGC TCCTAAAACC	16740
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GAAAGATAAA GAAATAAAC AGCTTCAGTA GGATCTGGAA AAATTTGATT TTATAGAGAA	17520
GCCTTTGTT ACAAACTCAA TATACTATCA ATAATAATAA TTATAGAAGC AACATAATT	17580
ATAATTCAC CTATCTGCAT CATTCTATTT CGAACTCTAA ATATATGTTC TATCAAAAT	17640
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CCAATATGCG TGACAGGTAA TAATGATAGC CAAAAATAGC AAGAGCAAGC AAGACGATAA	17940
GAGCTCCTAC TCCCAGCTG ATGGCAAGGA TAGGGGAGAG AGACTGAACC AAGAATATGC	18000
TCCCAATTAC AAGGCCATC AGGATTGCAC TATAATAAA CAATAAAACT ATGGCGACTA	18060
TGCCATTGAA ACGATTTCACC AGGTCCGTAA TGCTACTCCA ATTGGTTGAC AGATTTTAA	18120
CGTCCTTAAA GTAATGGTGG CAAGAAAGGA TGACACTGGC AATGATCCAG ACTACAAGAA	18180
GGTAAATCAT CGAAATGATG GGCAAGCCTA GATATAGAGA AAGACCAAGC AAAGTCGAA	18240
CTGGTAAAAA GGACTGGACA GCATATATAA TCCAAAATTT CACTTTCACAA TAACGAGCAA	18300
AGTCAAAGGG TAAACTCTTA AGAAAATCAA CATTTCCTCCT CTCCAAGGAC AAGGCAATTG	18360
AATGCAGGCT GGTGATATTG TTATTGACAA CTGCTATAAA GAGAGCTATA AAAACAAAGG	18420
GTAACCAGTA TGGAGGATGA ATGTCTGGAA CTATCTGAGA ATCTCGGATT TTGAAATCA	18480

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GACCGATCAT CATGAGATAA GGAAGGAAAG CACTTGTAAA AAGCACTGTA ATCACGCCAG	18540
TCCCCGTCC CAAGAGGGTG AGGTGGTAGC GTAAAACCAT GCGAAAAAAT CCCTTTTAG	18600
TGGTTGAAAT TCTCTCCTTG CTGCGACGTT CTTTTTGAC CTTCTCCTCA CTATTAAGCA	18660
GGATCACGTC ATAAAAACGA GGAAGGACCT TCTTTTGTT CAGATAAAGC AGGAAGAGAG	18720
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TTTTGTAAA GATATGAAGA GGATAAAAGGA GAAATGGAAT GTCTCTAACT TTGTCAACAA	18840
TACTTCCAAA AGTCGACTGA AGAAAAGAAGA TAAATATTAA AGGTATGAGA ACTCCTATCC	18900
CAATCATCAC ATTGAAAAA ATAGACTGAT ACTTTCTGAA GACCCTAGTT TGAGCCAAGA	18960
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GCATCAAAGG CAAACCCAGC CATAGAGAAG GAGCTAGCCT AATGTAGAGG ACCAGAAAAT	19080
AAGCTAGGAT TGGTACAATT CCAGTTAGAG CTGGCAAAAG GACAGACAGT CCTTTAGCAA	19140
TTATAATCTC TGATTCTTTA AAGGCATAGG GCCTATACGA TACCAAATCC TTACTCTCAT	19200
AAAAGACATT GTAAAAGGC GTAAAGAAG TTGAAAAGGC AATCACTAGT AAAATAGCAA	19260
TCATCGAGCT AAAATAATAA GGTATTTCT CAAAGGAAA ATGAATGGCT ATATTACTAA	19320
AACAGATGAT CATCAAGAGA CTGGAAAAAA TGTAAGAACT TAAGACTCTA GCGGAAACAT	19380
TTACTTTTT	19390

(2) INFORMATION FOR SEQ ID NO: 87:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 18436 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 87:

CCGAGCGTCG TTACAGACTT TATCAAGATT GGACGCAAGA AGAAATTCAA CATATAAAGG	60
AAAATATGGC ACAATCTCCA TGGCATACTC ATTACCATGT TGAGCCAAA ACAGGACTTC	120
TCAACGACCC AAATGGCTTT TCTTACTTTG ATGGCAAGTG GATCCTCTTT TACCAAAATT	180
TTCCTTTGG TGCAGCCCAC GGTTTAAAT CTTGGGCACA GCTAGAAAGT GATGATTTGA	240
TTCACTTTAA AGAAACTGGA ATCAAAGTTT TACCAAGATAC TCCATTAGAT AGCCACGGTG	300
CCTACTCTGG TTCTGCCATG CAATTTGGCG ATAACCTATT CCTATTTAT ACAGGAAATG	360
TTCGCGATAA AAACCTGGATC CGTCACCCAT ACCAGATCGG TGCTTTGATG GACAAGGAGG	420

GTAAGATTAC	AAAGATTGAC	AAGATCTTGA	TTGACCAGCC	AGCAGACTCT	ACTGACCACT	694	480
TCCGGCATCC	ACAAATTTT	AACTTCAGG	GTCAATATTA	TGCCATTGTC	GGCGGACAAG		540
ACTTGAGAA	AAAAGGTTTC	GTTCGCTCT	ACAAGGCTGT	CAAATAACGAC	TACACAAACT		600
GGCAAGCAGT	TGGCAGCCTT	GACTTTGCTA	ACGACCGTAC	TGCCTACATG	ATGGAATGTC		660
CTAATTGGT	CTTTGTAGAG	GAACAACCTG	TCCCTCTCTA	CTGTCCACAA	GGATTGGATA		720
AGAAAGTTCT	AGACTACGAT	AATATCTTTC	CAAATATGTA	TAAGATCGGG	GCTTCCTTG		780
ACCCCTAAAAA	TGCCAAATG	GTAGATGTGT	CTCAACTTCA	AAACATGGAT	TACGGTTTCG		840
AAGCCTATGC	AACTCAAGCC	TTCAACGCTC	CTGATGGCG	TGCTCTAGCA	GTTAGCTGGC		900
TTGGTTTGCC	AGATGTTCT	TACCCATCTG	ACCGTTTGA	CCACCAAGGA	ACCTTCTCTT		960
TGGTCAAGGA	ACTCACTATC	AAAGACGACA	AGCTCTACCA	GTATCCAGTC	GCTGCTATTA		1020
AGGACCTTCG	TGCTTCTGAA	GAAGCCTCT	CAAACCGTTC	CCAAACCCAAG	AAACACTTACG		1080
AACTTGAAC	CAACTTGGAA	GCTAAATAGCC	AGAGCGAGAT	TGTCTTACTT	GCTGATAAAG		1140
AAGGTAAGGG	ACTTCAATC	AACTTTGACC	TTGTAAACGG	TCAAGTAACA	GTGGATCGTA		1200
GCCAGGCTGG	AGAACAGTAT	GCCCAGAAAT	TTGGGACAAC	TCGTTCTTGC	CCTATCGAGA		1260
ATCAGGCTAC	TACTGCTACA	ATCTTCATCG	ATAACTCTGT	CTTTGAAATT	TTCATCAATA		1320
AAGGAGAAAA	AGTATTTCT	GGTCGTTCT	TCCCACATGC	GGACCAAAAT	GGTATCCTGA		1380
TTAAATCTGG	AAACCCAAC	GGAACCTTACT	ATGAATTAGA	TTATGGTCGC	AAAAACTAACT		1440
GATGTCGCCA	AACTTGCAGG	CGTCAGTCCT	ACTACCGTTT	CTCGGGTTAT	CAATAAAAAA		1500
GGGTATCTAT	CTGAGAAAAC	CATCCAAAAA	GTCAAATGAAG	CCATGCGAGA	ATTGGGCTAT		1560
AAACCCAACA	ACCTGGCTCG	TAGTCTGCAA	GGAAAATCAG	CTAAGTTAAT	CGGCTTGATT		1620
TTCCCCAATA	TTTCCAATGT	TTTCTATGCA	GAATTGATTG	ATAAATTGGA	ACACCAACTC		1680
TTCAAAAATG	GTTACAAGAC	CATCATCTGC	AAACAGTGAAC	ATGATTCTGA	GAAGGAACGC		1740
GAATACATCG	AAATGTTGGA	AGCCAATCAG	GTGGACGGCA	TCATTTCTGG	TAGTCACAAC		1800
CTAGGAATCG	AAAGACTACAA	TCGTGTGACA	GGCCCGATTA	TTTCCCTTGA	CCGAAACCTA		1860
TCGCCAGACA	TCCCTGTCGT	CTCCTCTGAC	AACTATGCTG	GTGGGGTTCT	TGCTGCCAA		1920
ACCTTGGTCA	AGACAGGTGC	CCAGTCTATC	ATCATGATTA	CAGGGAATGA	CAATTCTAA		1980
TCGCCAACCG	GACTGGCCA	CGCTGGTTTT	GCATCCGTAC	TCCCAAAAGC	TCCTATTATC		2040
AATGTTCCA	GTGACTTTTC	TCCCGTCAGA	AAAGAAATGG	AAATCAAGAA	TATCTTGACC		2100
CGGGAAAAAC	CAGATGCCAT	TTTGCTTCG	GATGATTGAA	CAGCTATTCT	GGTCATTAAA		2160
ATCGCTCAAG	AATTGGGCAT	TTCTGTCCCA	AAAGAGCTCA	AGGTCACTGG	CTATGATGGG		2220

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ACCTACTTTA TCGAAAATTA CTACCCCAA TTGGCTACTA TCAAGCAACC	2280
ATTGCTTGTC TCACATTGAA TCTTCTCTTG CAAAAGATTG AAGGCAAGGA AGTCGCCACA	2340
ACTGGTTACT TCTTACCAAGT TACGCTATTAA CCAGGAAAAA GTATTTAAC ACAAGAAAAC	2400
TCAGACCGAT TCGTCTGAGT TTTTATGATC TAAATTTC GAGATAGCGC TGGCTGTCT	2460
CTAGGTTAAA GGTTTATCT GAGATGAGGC GCTCTACTAG GGGAGCAACT TCAGATTCAC	2520
TAGCCCCAGC TAGGAGAGCT AGGGATTGG CCTGTAGTTT CATGTGGCCT TGCTGGATGC	2580
CCGTACTTAC CAAGGCTTTG AGGGCTGAA AATTTGAGC AAGACCGATG GACACGATAA	2640
TCTGGCTAA TTCTCTGGCA GAAGGATTTTC CTAGTAGATC ATGACTGAGA ACTACACGTG	2700
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CCAATTCTTC TCTTCAAGG TCCAGCGTCC AGCAGCTAAG ACCTTGATAG CGTCCATCTC	2820
GACTGGCAAA GGCATGGGCC CCAGCTTCGA TGGCACGCCA GTCATTACCA GTGGCAATCA	2880
AAATCCCATC AATACCAATTAA AAAATTCCCTT TATTATGAGT AGCAGCTCGG TAAGGATCAG	2940
CCTGCCAAA CTGACTAGCC AACGCAATT TCTCCGCAAT CTCTCGTCCT TGATCCTTTT	3000
GGCGGCTAA GTAGCGAAAG GCGATGCGAC AGCTTGCAGT CACCAAGAGAA TCGGTCGGCT	3060
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TATAAACAAAC GAGAAAGTCT GGTCGCCTT TTATCTGTC GACATGCAGA TCACCGGCC	3240
CACCTCCACG TTTAACGATA GAAGGATAGG CTTGATTGGC AACCTCCAAG AGCTCCGCTT	3300
TCTTGCCTGGC AATCTTCTCT TGCGCTAGTT TAGGATTAGC AACTTGATAA AGGGCTACCT	3360
GCCCAATCAT CTGTCGCTGA TGGACTTGTG CAGTAAACC ACCTGCACGC TTGATGATT	3420
TGCTGGCATA GCTGGCCGCC GCAACCACAG AGGGTTCTTC TGTCACATAG GGAACGGTGT	3480
ATTCCTGACC GTTGACAAGT ACCTCCGAA CCAGTGAATA AGGCAGAGAA AAAGTTCCC	3540
CTACATTCTC ACTCAGCTGG TCTGCCACAG TCACGCTCAT CTGTTCATCC TTCTCCAGAC	3600
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ATGATTTTT AGAAAATCCA TTCCAACCTA TCTTCATTAT TTTTCAACCT TGCTATAACG	3720
GGGTTGGTGG TCGAGAATT CAACCAAGGC AAAATCTGA TTTTCATAGC CAGCAAAC	3780
GGCAGAGTTA GTTCATCCA AGTTTACTTC CTCAAAAAG ACCTTTCAT AGTCTGCAAC	3840
GGATAGGGCA GTTCGTTGGT TGAGCTTGT CAAACGGTCT TTATCCAAT AAGCTTCATA	3900
TCCTTCAACC AATTCAACCAAC TGAAGAACTC AGCCACAGCT CCACCTCCGT AACTATAAAG	3960

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GGCGATTTA TCCCCAGCTT TCAAGCTATC TGATTTTCC AAGAGAGACA AAAGTCCAAG	4020
GAAAAGTGAA CCTGTGAGA TATTCCCCAC CTTTTGACTG TAGAGAACAG ACTGGTCAA	4080
ATGCTTTGT AAGAGGTCTT TTTTCTCTTG AGGCAGGCTC TTATCCATGA TTTTTTCAA	4140
GCCTTTAGC GCTAATTAG GATAAGGCAA GTGGAAACAA ACAGCCGCAA AATCATCAA	4200
AGTAAGCTGG TAGCGTTTTT GATATTCAAG CCAAGTCGTT TTCAAACATAT CCAAGTATTG	4260
TTGGTAGAA TAGACACCAT TTACATAAGG AGTTGTGGAG TAATTGGTC GCCAGAAATC	4320
CATGATGTCA CGGGTCTGAG CTACATTGTC ATTATTAAG GCCATCATGC GTGGATTTG	4380
TGTAATCAAC ATAGCTACAC TTCCAGCACC TTGAGTTGGT TCTCCTGGAG TTTCAATACC	4440
GTATTGGCA ATATCACTGG CAATGACCAA GACCTTGGAC TCCGGAGAAT TTTCCACATG	4500
CAATTGGCA TAATGGAGGG CAGCAGTCGC TCCGTAGCAG GCTTCTTAA TCTCGAAACT	4560
ACGAGCAAAG GGCTGGATGC CCAGCAAGCC ATGCACAAAG ACAGCCGCGAG CCTTACTCTG	4620
GTCAATTCC GACTCGGTG CCACAATGAC CATGTCACT TCTTGTCTTT CTTGCTCAGT	4680
TAAAATAGAG TCACTAGCAC TGGCCGCCAA GGTCACGATA TCCTCAGTTA GGGCGCAAT	4740
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CACAAATGGC AGTAAAAGAG AGAAAAAAAGA CTTGATTCAAC CAAATCAAGC CTCTTATTGG	4980
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CTCCATTCAAC AGTTATCAGC TCCTGGAGGA TCAAATTTCAG TCAGTAAAGTC CTTCCCACATCT	5100
CATCTACAAA TTTTGATAA AACTGACTGG TCGGAATTTC TCTGACATCC TTATCAAATG	5160
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GAAACTCTGA GCCCGAACTA GAACCATGA CTGGGATAAA CAACAAAGTC AGTAGATTTA	5280
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ATTTTACAG CTTGTCGAT ATTTCTGCAA CTCATTTGA ACAACTTCAT ATCATGACTA	5640
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ATTTTTACTT CCGTCACATC AATCATTATC GTGTCCCAA AGCTGAGAGG ACTTCTTGAA	5820
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CTACCGATACT CGATGTGTTT TTATATAATG ATAGAGTCTG AGAACACTG TTCCGCTAGC	6060
CATTCCAATA GAGATTACCA AAGCCAACAT GACAACCAAG GTCGCACTTG CCAGTGCTTT	6120
ATTATAGTCC CCTGTACAAA AAAAGGCAGT TGTTCGGTAG GAGAGATAAC CTGGAACCAG	6180
CGGTGCCAAA ATGGCAAGA TAAAGACAC ACCAGGTGTC TTATAAAGAA TACTTAAAT	6240
CTGGCTGACA CAAGAACCA TAATGGCTGC AATGAAGGTA GCTACAAATGA CATTGGTCCG	6300
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CATAGACCGT TGACACATTGA GTACGATTAA AAAAGTCATA ATGGCAAGAA AACTTGCTAC	6420
TGCTTGTAAAT AAAAAGGTTG TTAGTGTCAT ATTAGTTCAT CAATACCAAG GCGACAGAAG	6480
TTCCCTGCCCT TAAAGCGAGG GTAATGAGCA GGGATTCAA CATCTTACTC ATACCAGAGT	6540
TTATGTGGTT GGTCTATAATA TCACGGACCG CATTGGTCAA GGCAATACCT GGTACAAACG	6600
GCATGACCGC ACCAGCTATA ATCAAATCTG CCGTTGAAGG AAAACCTGTG TAGCGAGCCC	6660
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TCGACCAGTT CCAGCGCCAG GATCTGTACG GATAATCCGC CCTCTCTCCA CCTTTCACT	8100
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CACTGCCACT TCTTCCCCAT CTAAGATTAAGT GTCTTGGCT AGGTAGACAT CCGCCATACC	9240
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ACAATATCAC GAATCTCACT GCCTGAAATC ATGTTGGTCA AGCCGTCACT ATTGAGCAAG	9480
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TGACCAGCCT TGAGCAATTTC ATTAACCAAG GAATGATCGC TCGTCAACTG ATGGTATTCT	9660
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700	
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CTTCGATAAA GTGGGATAGG TACCAATTCCA GAGTCAGTTT ACGGGCTACC GTTCCATAGA	11220
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ACTTGACCTG CCTTTTCAGC CCGACCTGAT CGACCTGCCA CCTGAGTCAA GAGCTGGAAG	12780
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CCCCCAACC TTAAATTTT TCACCATTTT CTTTTCTTT AGCAATTGCG TCTTTGATTT	14820
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GTTTCTTC TGGAATCTGGG TGAAATTGAA CGTTCTGA TTCGATTCT TCTAAAGCGC	14940
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CATCTCCGTG GTGACTAGCA ATCGTATTCA CCACAACTGG GGGTTCTTG TACTTACGGG	16260
CCAATTCCAT ACCGATTTCAC CGGTGGCTAC CTTCAACCTC ATGGTCAATG GCTTTCCCGA	16320
TATCGTGAAG GAATCCAGCA CGACGGCAA GAGCCGCATT TTCACCAAGT TCGCTCGCCA	16380

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TGATACCAGC CAACTTAGCA ACCTCAATCG AATGGCGCAA AACATTTGT CCATATGAAG	16440
TACGGAACTG CAAACGTCCC ATAATCTTCA TCAAGTCTGG ATGAAGGTTT GGCGCACCAA	16500
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AGGACTACAA TGTCCCTCATC TGGATTTTA TTGACCACCG TCAAAGGCTC AAATTCATCA	18120

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TTTACCTCT TCATGTAGTC CTTAAATGA TTTCGGAATG TTGAGTAAAG GACTGCTTCC	18180
ATAACCACAC CTCGTTTAG CTCTTTCCA CTATTATACA CGAAAAGAAA GAAATTGTCA	18240
GGAACCTGTA CAAGATTTTC TTTCTATCT ATTTATACTC AATGAAAATC AAAGAGCAA	18300
CTAGGAAACT AGCCGCAGGC TGACTTGAG TACGGCAAGG CGACGTTGAC GCGATTTGAA	18360
TTTGATTTTC GAAGAGTATT ATTGTAAAAA AATCTCAAAA AGCCTACCTT TCGGTAGACT	18420
TAGTTGTTT CTATT	18436

(2) INFORMATION FOR SEQ ID NO: 88:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 7001 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 88:

ACGTAGAAAA ACTATTTCTA TCACAGATAA TATTCCGTAT CTTGTTGGAG GTATTGAAAT	60
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AGAAATTTG AATGAAGATT ATCCCTTTTC AGAAATCAAT TCCACCATTAA CCAATATGAT	180
AAAAACAGCT ATAGAGTACG TCCCTTTGGA AACAAAATTA CTTGGATTTG CCTTATCAAT	240
ACCTGGACAT TATAACAAAG ACTCCGGAAG TATCATTACA AACAAACCCCA TATGGGAATC	300
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ACACGCCCAA TTATTTATTA AAAATCCCA ACTAACTGTA CTAAAAAGCC TTGTAAAGAC	660
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CATCAATCCT GCTGATAAAA TCTATATCAA CAGTCATTG CTTAATTATC AACCTTTCAC	840
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TATCGTCGCT TTTTCATAG AACATAGCAA TGTTTACAA GATATTATTT CACCTTAATA	1020
TATTAGAAAT CTATAGACCT GTTAAATCA ACTATAACCT GTAGTAGATA TCTCGTATTT	1080

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AGACAAATATG AAAACAAGAC GACTTCCATA TAGGAAACCG CCTTCTCGCT ATGTTGAGTG	1140
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AACCCCCAGA ACTTAAATAA CAATTTTAT TCAAGATACA TACTCCTAGA ATAAAACTTA	1260
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AACAATCAAT TGATGCTAA ACCTGTACCT AGATGTTCG GTTCATAAAA CCATGAAACT	1500
GTAAAAGTGG ATGAAATTGA TAGCGATAGT CAAATCAAGA GGCATCATAA CTCTAAAAAG	1560
TCACAATATA TAAGTTCATC CTCGGAAAAA TATCATTCTA ATTGTTGAAA TGCCCTACATG	1620
AAAAGAAAACG TCAAATGCTC ATGAAACAAC GAATACAGGT ATCAAAACTA TGACAAAACA	1680
AATCCCTAAA TTTACTAAAG ACACTGCTCA ACTTTACACC TGAAATGGT TGTTGTATAA	1740
TAAAGTTACA AAGATGTACG ACCACACTGT TGAAATCAT AGTGTTCGCG AATATATTAC	1800
TGATAGCATT TCTACAAATA CAAGTAAAGA GAGCGGATGA GATTCAAACG AAATATGTCA	1860
GTGCTTTGGC ATTCCCTAGCC TTCATATCAT TTAAAGAATT CTATAGACAA AATTTTTTCC	1920
AATACAGACA CTCGTAACAA CTGCTTCATT TTTCTACCAA CATATTTAGG AACAGGATAA	1980
GATACAAAGAG TATTAATCCA TAGCTCAGTT CTATACCAAT CTAAGACAAA TAAGCTAAA	2040
AAACGATTGA TAATAAGCAA ATAGATTCCA AATTTTCTCT ATCTGCTCAT TTAAATAAAC	2100
AATACTAGTG TAACTATCCT TCCAGTCAGA AGCTTGTCAA ATCACACCGA AAATTCTTCT	2160
AAAATTTATC TCGTTAGGCA ATCAAGCAA AACTCGACGA TAGTACAAAC ATTATCATA	2220
AGGATTGACT TCCTAAATTAA TATACTTAG TAAGGTTTC GGATAAGAAA AAAGGTTCAT	2280
TTTACATTTC TAAACATTCT TTTCTAAGAT GAAAACAGA ATTTTCGAT TGTGATTAA	2340
AGCAACAAGA AGATTTTCAG TATCATCCTA TAGATACGAG CTAATTAAGA AAAACTACAT	2400
TTTTGAATAT AAACTACAAT AATATAAACT AAATTTTATA GGAGGAAGAC AATGGATTGG	2460
TACGATTATA TGATACAGGC ATCCAAACAA TCACAATTCA ACGCAAGCCA TTGGTTTCGC	2520
TATTTGCGAA AAGTTATTT TGAAGACTAT TCTTATTTAA CAAACCAAGA TGTAGAAAAG	2580
TTGCTAGACT CCAAAGAACT AACCCGTTT CAAAAAATTA GCTTGAAGTA TGCCCTTCAA	2640
GAGCATACTC CAACTCATAA ATATGTGATT TCATTAATAA AACCTGCTAA GTTAACCAAT	2700
GTTCAAAAAT TGATGGAGAA ATACAAACAT GGATAAAATG AAACCGGTCT TCCAAGGCCCT	2760
AAATAAGGAA TTAATTCAAGG AAAATCTGAC TTAAACAATT ATCTGTGTCG GTGGTTATGT	2820

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CTTAGAATAT CATGGTTAC GTGCCACACA AGATGTTGAT GCTTTATGG CTCTATAATA	2880
TTTGTAGTGG GTAAATCCCC TATGGATATT ATGGAGCCTA TTTTGTTGTA GAAAAAAAAGT	2940
CCCATATGAC CTATAATGAA AAGGCACAAA ACAACTCATT AGAAAAGAATC ATATGGAACA	3000
ATTACATTTT ATCACAAAAT TACTAGACAT TAAAGACCT AATATCCAGA TTTTAGACAT	3060
CATCAATAAG GATACACACA AGGAAATCAT CGCCAAACTG GACTACGACG CCCCATCTG	3120
CCCTGAGTGC GGAAACCAAT TGAGAAAATA TGACTTTCAA AAACCGCTA AGATCCCTTA	3180
CCCTCGAAACA ACTGGTATGC CTTCTAGAAT TCTCCTTAGA AAACGCGTT TCAAGTGCTA	3240
TCACTGTTCA AAAATGATGG TCGCTGAAAC TTCTATCGTC AAGAAGAATC ATCAAATTCC	3300
TCGTATTATC AACCAAAAAA TTGCGAAAAA GTTGATTGAG AAGATTTCTA TGACCGATAT	3360
TGCTCATCAG CTGGCCATT CAACCTCAAC TGTCATTGCG AAGCTCAATG ATTCTCACTT	3420
TGAGCATGAT TTTTCGCGTC TTCCTGAGAT TATGTCCTGG GACGTTGAAA CAGTCGGGG	3480
AGTGACTGTT TCAATCGGGA GATGGAGATG AGCTTTATTG CGCAAGATT TGAAAAGCTC	3540
GATATCATCA CTGTTCTTGA AGGTAGAACAA CAAGCTGTCA TCCGAGATCA CTTTCTTAAA	3600
TATGATAGAG CCGTCGGATG TCGCGTCAA ATTATTACTA TGGATATGTT TAGTCCTTAC	3660
TATGACTTAG CTAGACAATC TTTCCCGTGT GCTAAATCG TTCTTGATCG CTTTCACATT	3720
GTACAACATC TTAGCCGTGC TATGAGTCGT GTGCGTGTCC AAATCATGAA TCAGTTTCAT	3780
CGAAAATCCC ATGAATACAA GGCTATCAAG CGCTACTGGA AACTCATTCA ACAGGATAGC	3840
CGTAAACTCA GCGATAAACAA TTTTATCGC CCTACTTTTC GTATGCATT AACCAATAAA	3900
GAGATTTAG ACAACCTTTT GAGCTTATTCA CAAGACTTGA AACATCACTA TCAGCTCTAT	3960
CAACTCTTGC TGTTTCACTT TCAGAATAAG GAACCGGAGA AATTTTCGA ACTTATCGAG	4020
GACAATCTTA AGCAGGTTCA CCCTTATTTT CAGACTGTCT TTAAAACCTT CCTCAAAGAT	4080
AAAGAAAAGG TTATCAACGC CCTTCAACTA CACTATTCTA ATGCCAAACT GGAAGCGACC	4140
AATAATCTCA TCAAACCTTAT CAAGCGCAAT GCCTTTGGTT TTGAAACCTT TGAAAACCTC	4200
AAAAAACGGA TTTTATCGC TCTGAATATC AAAAAGAAA GGACAAAATT TGTCCTTCT	4260
CGAGCTTAGC TTTTTTCAA CCCACTACAG TTGACAAAGA GCCGGAAAAA GGAACAGCCT	4320
TAGCTTCCCT TTCATTCTT TTTATTTCCC TCGTAGTAA CGTGTAGCT TCCACAAAAC	4380
AAACAGGATT CCCAGAAATG CCAGTACACAC TAGCCCCACGG TACAACCATT GAGAGGTTGC	4440
AACACCGCAT ACAGATTGTC CTTCTTTCGT AAAAGCAACC CTCGCAACTG CAGCTGTTG	4500
TGGATCTGAT TTTTGATAAA CAGCGACTCG TTCAAAATTC ACTAATAAGC GTTTATTAAA	4560
GGTAGGAATC GGATCGCAGG TTATCAAGGT CATGATATT TTAGAGCTAA CCGATTCTAA	4620

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TTTTTCCCAT TCCGACGGTA AAATAATCTC TGTGTCCATC ATCTGATATT CTACAATTTC	4680
CTGGCCATTA TCATAATAAA GAGCATCTCC AACCTTTAGC TGATCCAAT GGCGGAAAAA	4740
GACATGGCTT GGCTCTGCAC GGTGCCAGC AATCACTGAG CGAATCCCTG TACCATCCAG	4800
AGGCAGCGGT GTACCATCCA CATGAGCAA GCCCATCCCT AAATGATGAT AATCTGCTCC	4860
CAAATAAACC GGCTCCATGA TTCCAAAATC TGGAATGAC AAGTAACCAT AGACTGCATC	4920
AGGGTCGTCA GACACTTGGT AATTGACCTC ATATCCCTCC GCCAAAAAAG GATCTACAAT	4980
GCGATTTCGC GAAGCCAAGC GTTGATTGTA GCGGAGAGAA TGGTTCTGTT GTTCTTGGTA	5040
CATTTCAAGT GTCACTGGATT TCACAAATGT AGCATGACCT TTCACCTGTC CAAGAGACTG	5100
CAACACCATC TGTCCAAAAC AATAAAATAGG AATCAACAG CGTACCAACA TCAACAAAGTA	5160
TCCCAATAAG GCTCGTAGTT TAGCCTTGA CATGACGCC CTCACATTGC TTTCTAGTC	5220
CTTTGACAAT CCGTCGATTA CGATACACGC GATACAGCAA GAGAAGGATG ACCGCCATCG	5280
CTCCTAGTAA TAACCACAAAC CAGAATTGCC CACGCTCTCT CACCGCTCGA TTCCGCTCTG	5340
CAATTGGTGC CGTATACGGA ATCCGCTTCC CACGTACCAA CAGACGATGA CTGTTAATCA	5400
TATACGGTGT ACAAGTCAAC AAGGTCGCAT AATCTTCCCC ATGTTGAATC AAGACAGGCT	5460
CAAAGTCATT CGGCTCCACC GTCACTATCT GATCCACTTG GTAGGCCAAC ACCTGATCTA	5520
AAACGTGAAG ATAAAAGATA TCCCCTTTTT TCATCTTATC CAATTGACTG AACAAATTCTG	5580
CCGTGGCAA TCCTCTGTGA GCAGTGATCA CTGTATGGGT ATTTTCACCT CCAACAGGCA	5640
GCGAAGCCCC TTCTAACAGC CCTGCCCTT TCTGAAGAAT GTCCTCACTC GTTCCGACAT	5700
ACATCGGAAT TTCTGTATCA ATCCGAGGA TTTCACATA GCCAACCGC TCATGGACCT	5760
TTAGCATATT GGCACTATTCT GAGACGCCCTT TCTTTTTCTC TTGCTCTGTA AAAGGATCAA	5820
GAATTTCAAG TGGTTCAAG GTCGCATTGA AGGCTTGAGC CAAGGCCAA CGCTCCTCAA	5880
GTTCTGCCCTT ATCCATCTGG GAAACCGTCT CATCAAACTC TTTAATAACC TCGTTTGACT	5940
CAATACGATA ATAATAACGA GACACCAATG GATATATCGC AACGGCGAAT CCTACTAAGA	6000
AAATCAGAAG AAGGATCAGC GGATGTTCTC TCTTTTTGT GCCTTTTTT CGTGAACGTC	6060
TACTGTTGTC CATCCTCCAC CTTCACTTCC TTCCCTGCTG CTTTCAGCGC CTTCAAAGCC	6120
TTTTCGGTT GTTTTTCTT CTTGCGCAAG CGTCGAATAA TCCATAAAAG AATCACAATC	6180
AAACCAACTG CCACATAAAA CAGGTAGCGA TAGAGATGAC TGAGTTGTT TGCTGCAATA	6240
AATTCTTCCCT CAACCTCTGC TACGTACGGT ATCCGATGCC CCCGAACCAA TAGACGATGG	6300
GTATTGATCA TGTATGGCGT ACAAGTCAGC AAGGTCACAT AATCATGACC TGGTACAATC	6360

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AATAAAATCAT CAAAGTTCGT CGGCTCAATC ACCTTTACTT GATCCACTTG ATAGGCCATC	6420
ACTTCCTTGA TATTGTGCAC ATAAAACCTTA TCCCCAACTT TAAGTTGGT CAAATCCGTA	6480
AACATCTTAG CTGTTGGCAA ACCTGTATGT GCCGTAATCA CCGCATGGGT CGAATTGCC	6540
CCGATCGGCA GAGAAGTTCC CTCTAGATGC CCAGCCCCTT GCTGCAATAC CTCTTCAGCA	6600
GTACCAGCAT AAACCGGCAA ATCCACGTCA ATAACGGGGA TTTCACATG CCCCACATCCGC	6660
TCATGGATT CTAAACATACG TGCAACTCT GCTCGCCCTT TTTTCTTCAT TTCTTCCGAC	6720
CAAGGATCGC CACTCACTAC ATTATTCAA GAGTCATTGA AGGCTTGTGC CAATTTCATT	6780
CGTTCATCAA TGTCAGCCTC ATCCAACGTT GCTTTTCCT TATCAAAGTC AGCAATTGT	6840
TGATTGATT CCACTCGATA ATACAAGCGA GACACCAGCG GATAACCCAT TACCGCCATT	6900
CCAATGAAAA ATACCACTCC TAATAGGAGA TTATTCGTT TTTGCTTTTT TGTTTTTACC	6960
ATTTTATCA GCATCCCTTT ATCTTCAAAC TTCAGGGTAT C	7001

(2) INFORMATION FOR SEQ ID NO: 89:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 10411 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 89:

GAGGGAGCTT AAGAAGTTAC CACCGTCCTC TAGGCCCTTA TCCGCATCAA AGTTAAGGTT	60
GATATTTTTA AAACTGTGCGC CAGCTTGTGA TACGATGCTT TGTAAAGGT CATTTAGGGT	120
TTTAGTGAAA TCTGCATTGC TGAGGATATC ACTCTTTGAG AGATTCAAGG CAAAATTGAT	180
GATGATATTG ATCTGGTTTC CTGTTATGAC CTGATCAAGT TTGTAATTTC TTAAGGTATC	240
TTCAACAATC TTGGGGATAT CTCTCTCTGT CAGATTCCC TTACTTTCTT TAGCTTGGC	300
GAGTCCTGAC TTGATATCAG CTAGGGCAC CTTAAATTAA TTAGCATCAT AGCCTGATTT	360
GTCCTTGTCTT TCAGCATTGA TATCTGACAA AGCTTTAGC TCTTCTGAG CCAAATCTT	420
ATTAGCTTGT GGCACTTGG CTCCATTAGC CTCTAGCGAA TAGTAAATCC CTGCTAAAGC	480
ACTTTCTCCT GTAACTGGAA TAGGGGCTGC TACAGTGATT TTGGCATGTT CCATACCCAG	540
CGTTACTGCT GCGTTTCGGT ACATATCCTG AGTCACCTTA GTGATATTTC CTGGTGTTC	600
AATCTTGACC TCAAGTGGCG ATTTGTCAAC TAGCTTTGA ATCTTGGCTG ATGAATACAA	660
CTGTAAGCTA GAGTCATTGG CCACATTCAT GATTTAGAA TAAACATCAG GTGTCATGGT	720
CTTGAGTTCT TTGGTATCTG TTGAGGCATT GTAGCCCAGT TTTTAAGAG TTTGATTTT	780

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TTGGTCTTCA	GATAGGGAGG	AACCTAGGAC	ATATTCAAGGT	TGGACATAGG	TTTCATCGAT	840
AACTTTTGTA	ACATCTGTTG	CTGCATGGAC	GCTATTCTA	GCTGTTACTG	CCACACAAGAT	900
CGCAGCGCTA	GTCAGAAAAGA	GTTCCTTCT	CATAGGGAA	TTCCCTCTT	ACTTCTTTAG	960
AGTAATATAT	CTATCTTAA	GAAAACTTAT	AACAAAAACA	CCTGGTCTAG	CCAGATGTTG	1020
AAAAGAGAGT	GAAACATTG	ATGATGTTAA	GGTTAACGTC	TACCTGTCTA	GAATAATAAT	1080
AGTTCCCTCC	ATTTACATAG	AGTCAGCAC	CGTGAAAAT	GGAAATGGGG	TGAATATAAC	1140
TATAAGTCTT	TCCAGTCCTA	TTACCAAGCA	AGGGGGCAAC	AGTCTCACGA	GAGTACTGTT	1200
TGGCTAGAGC	CAGGGTATTT	TCCTGCCAT	TTTGGGGAT	AAAATCGATA	TAGGCAGGTC	1260
CAAAATTATA	GGCTTGAACA	GCTGTCAGA	TATCTACCCC	CTTCTTCTGC	GCCAGATAGA	1320
GATTGCCCTGT	CAGAGTTGTA	ATGCCTTGCC	GAATGCTAGA	GGCATTTATCA	TTGATGGTGT	1380
TGGTGGAAACC	ACTTGAGAC	TCACTAGACT	GCATAACATC	GCCTCTTTT	CCTTTTGT	1440
CAGTATAAAT	CATAGCAAGC	ACAAGCTCTT	CGTTTGCTGG	GGTGTCTTGT	TCACTCAATA	1500
TTTCTCGCAC	CATGGGTTGTA	TAGGTCATGA	CTTGTGTTGAC	ATCTTGATGA	ACGGCGTAAG	1560
CTTTATAGCC	AGCAAAAAGG	AAGACTGCTA	GTACAAGCAC	TCTTCGAATT	CGTTTAAACA	1620
TTATTTACTT	TGGATATCCT	CGATATTTTT	GATTAAGATA	GAGTAGGTT	CATTTTCGTT	1680
TTGGATAAAC	TCAACAGACT	CGGGCTCTTG	ATAGACGTTA	TTGGGAACGA	TGAGCTCAAT	1740
TCCATTGAT	AAGGAGAGTT	TTGGTTTTC	AAATTTCTTT	AATTGGCGAC	TGGCATCAAT	1800
TTCATCAAAT	TGAACAGGTT	CTGGTACGGC	TTCTTGACT	TGGTCAATAA	AGCTCAAACG	1860
AGCCGTCAGA	TTGTTGTCAA	AAAGGTATT	AGCCAATTTC	TCAGGTGACA	ATTCAATTGCT	1920
TTCTCTAGG	TTGTTGAAA	TAGCTGATT	GACCTTGAT	TGAAATTGAA	AATCATCTGT	1980
GTAAAAAGAT	TTGCAATT	TCTGGCTGT	TTTTCCAGT	TCCTTGATAG	ATTTTTAGG	2040
AGAAAATCTTA	GGAGCGACAG	CAAGAAGATT	ATCTGAAAAA	TAGTTCAAAA	AAGTCCCCTT	2100
GTACTTGATT	CGTTTTCAA	TCAGGTGATA	CTTGCTACTC	TGAAGATTGA	CCACCAAGGC	2160
CTCATCAGCT	CCTGTTCAA	ATCCAGGCAG	GTTATTCTGA	GTTAGCTTGA	TTGGATTATC	2220
AACTTCTCCT	CCGAGGTGGG	TCAAGGTCTC	CCGCAGGGCA	ATTCGCAAGA	AAGCGAAATG	2280
TTCTACACCT	TCTTAGAAA	ATTGCACAAA	AATCAAGTCA	TTGGTCTTGA	GATTTTCAGA	2340
AATGCTAAAC	TCCTCTTCC	AGAGATTAGC	CAGCGTTACT	GATGTCTCCA	ACAAATCGTC	2400
TGTAATATGA	TTGAGAAGG	GATTTCTTC	TTGAAAATC	CCAGTCTTGG	CTTCATCTGA	2460
ATACACATGT	TCAATTTTT	TACCGAGGTA	TTCTTCGATT	TTGGAGTAA	TATTGAGAAA	2520

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CTTATCTGCT AAGAACAGTT CGGTATCATC CGGACTGAAC TGGTGAATAA TGGCTTTCTT	2580
AATATAAAATG TCCATAAAAG TTTTAGTCCT CGTATAATGG GAAGGCATCT GTCAATTCTT	2640
TGACTGCACT TCTCACTTCT TCTAATACAG CCTCATTTC TGAAATTCTTA AGGGTTTTAA	2700
TGATGAGTTC AGCCACTTTG CGACTTTCTT CTTCACCAAA TCCACGTGCA GTAATGGCTG	2760
CTGCTCCGAT ACGAATCCA CTTGTCTTGA ATGGTGACAA GCTTCGTAA GGGATTGAGT	2820
TTTTATTAA GGTAATATTG ACTTCATCCA ACAAGTTTG AGCAACTTTG CGCTTTCTA	2880
CAACTTTAGT CACATCAACA AGGAAGAGAT GGTTTCAGT TCCACCTGAA ATAATACGGA	2940
AATCAGGGTC TTGCAAGAAG ACATCTGCCA TAGCCTTGC TGTTCTTAATT ACATTGGCAG	3000
CATATTCTT GAAGGCTGGA TCCAAAACCTT CTTTGAAGGA AACTGCCTTA GCCGCCACAA	3060
CATGCTCTAA AGGACCGCCC TGAATACCTG GGAAATAGC TGAAATTGATT TTTTTAGCAA	3120
GTTCTTCGTC ATTGGTCAA ATCAAACAC CACGAGGTCC ACGAAGGGTT TTGTGGGTG	3180
TTGTTGTTGT GATATGAGCG TATGGAACTG GGCTTGGATG AAGGCCAGCC GCAACCAAGC	3240
CAGCGATATG GGCCATGTCC ACCATGAGCT TCGCACCGAC AGCATCTGCG ATTTCACGGA	3300
ATTTTGAAGA ATCGATAATT TGAGAATAGG CTGAAGCACC AGCTACAATC AGTTTTGGTT	3360
TTACTTCTTG GGCTTGTTC AAGATAGCAT CAAAGTCTAA GAGTTCCGTT TTAGGATCAA	3420
CACTATAAGA AACAAAGTTG TAGGTTTGAC CAGAGAAGCT AACAGGAGCC CCATGAGTCA	3480
AATGACCACC TGATGCCAA TCCATTCCCA TAACCGTATC ACCTGGCTCA ATCAAGGACA	3540
TGTAAGCCGC ACAGTTAGCT TGGCTTCTG AATGTGGTT AACATTGGCA ATTTAGCAC	3600
CGAAAAATTTC TTTTGCCTG TCAATAGCAA GAGTCTCTAC AACGTCTACT ACATCAGTTC	3660
CACCATATAA ACGGCGTCCG GGGTAACCCCT CGGCATATTG ATTTGTCAAG ATAGACCCCTT	3720
GAGCTGCCAT AACAGCCTTG GAAACTACGT TTTCCGAAGC AATTAACCTG ATATTATTTT	3780
GTTGGCGTTC TTCTTCTTGT GCAATAGCAT TCCAGAGATC AGCATCATAT GCTTTAAAT	3840
CATCTTGTC AAAAATCATA GGTCTTCTCC TTATTTGTGT GACTAGTCCA TTAGTTGAT	3900
TTTACAATAA GAAAATCAA CTAACAGATG CGAATAAACC GTTCTGCAT TTTATCACAA	3960
GTATAGCCAA CTTTTTCATA AAATGCATGA GCACCCAGAC GATGATTGGC AGAATTAAAG	4020
CGGATAAAACC CATAACCACA TCTTTTGCT TCTTCTTCCA ACCCTTGTAG TAAACTTTTA	4080
CCAATACCTT GACCTTGCAC TTGAGGTGAA ACTGCTAAAG CTAAGATATT AAATCCTGCT	4140
TTGGAATAGA GTGATTCGTA AACTTCAGCG TGGACATATC CAAGTAAGAC ATGATTAGCT	4200
GCATCCTCAT AGCCAAGTAG GAAATGATGG GAATCCTGAG ACAGTCTAGC TAGTTGGCTA	4260
GCCGTTCCCT CTGGACTAAA AGTATAACCC AAAGCCTCTT GGTTGATGTC ACATATAGCT	4320

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TTCACATCAG	TTTCTCTTAA	ATCTCTTAGC	ATCTCATTC	TCCTCAAAAG	AAATCTTGG	4380
CAACCGAGCA	AGAATATCTT	CTCGCTTAAT	GGCCCCTTGA	CGTAAGATT	TCACCTTGTC	4440
TCCCGACAAA	TTCAAAATAG	TTGAATCCTG	TCCAGTTAGA	AAAGCATCGT	CTTCCAGACC	4500
CAGAACCTCT	TGGTCAAAT	CCTCTAGAAT	TTGATTAAAG	GTCACTCCAC	TCGCCTGACC	4560
TGAGATATTG	GCAGACGGCC	CAATCAAGGG	ACCTGTCTCT	CGAATCAAAT	CAAGGGTAAT	4620
GGGATGACTA	GGCATCCGAA	ATCCAACAGT	TGCAAGGCCA	GAATTGACCC	AATAGGGAAC	4680
TCGGTCATTA	GCTTCGAGAA	TAATGGTCAA	GGGACCTGGT	AAAAAGATCT	CTACAAGTTT	4740
TTGAAGATAA	GTTGGCTGAT	TCTTGAAAAA	GTACAAGATG	TCCTCTAAAG	AGGCAACATT	4800
GAGATTGAGC	GCCTTGTCTC	TACGTCGACG	TTTAAGCTGG	TAAACATGGT	CAACTGCTTT	4860
TTCGTCTAGC	GCCTTAGCAA	AGAGACCGTA	AACTGTCTCT	GTAGGCCAAA	CGACAGCTCC	4920
ACCATTTTCC	AACTCTTGTC	TAATCCTGTC	CATCATCAAC	GACAACCAC	CTATCTTGAC	4980
CAAATTGGTC	CTTGAGTGTT	CGTACTCGCT	TTTCAGGAAG	ATGTTTCCTA	AAAAGTTCA	5040
GAACACTTTG	ACCTTGCTTG	TATCCAATT	CAAGGTAAAT	CTTACACCACCA	TCTTGAGAT	5100
AGTCTTTGTC	ATCTCCGCA	ATTCTACGGT	AAATAGCTAG	GCCATCCTCA	TCTGCAAAGA	5160
GAGCTAGATG	AGGCTCCGAA	TACAAGACAT	TCAAGCCTAC	CTCTGACTCA	TCTTCACGAG	5220
AGATATAGGG	TGGATTGGAA	ACAATTATAT	CATATTTTC	AGAAATTTCT	GTAAAACAGT	5280
CAGATTTTTT	AAAAATATT	TGAAGATTTT	GATTTTTAGC	ATTTTCGCTA	GCTACATCTA	5340
AAGCATCTTG	GGAAATATCT	GCTGCCGTCA	CTGACCAATC	TGGTCTGTTT	TTTGCTAGAG	5400
CGAGAGCAAT	AGCTCCACTA	CCCTGTCGGA	TATCTAGGAC	CATAAGATT	TTCACAGGAT	5460
TTTCAGCCAG	GATAAGCTCC	ACCAACTCCT	CTGTTCTGG	ACGAGGAATC	AAAACCGTT	5520
CATCCACCTT	TAATGCATT	CCATAAAAAT	CTGCCTGTCC	AATGATGTAC	TGAGCTGGCT	5580
TGTGAGCTGC	TAGTGCTGG	AAAATATCTT	CTACAAATTG	TTTTCTTCTCC	TCTGTTGTCA	5640
CCTCCTGCTG	GAGGGCAAAA	ATAAAAGTCTG	AAAAGATAG	ATTTTTCTAGA	CTACGATAGA	5700
CAAAAGAGAG	GCTTTCCGCT	TCCTCTCCTT	GTCTTATCAA	CTCTTCTTCA	AAATTTGAAA	5760
ATAATTGAGC	TAATTCATT	ATTGTTAA	TTCTCTAGT	TTTTGTGTTT	GGTCATAAAG	5820
CACCAAGGCA	TCCACAACTT	CGTCCAATT	ACCAGACAAA	ATCGTATCTA	GTTTTTGGAG	5880
GGTCAAGCCG	ATACGGTGGT	CTGTGACACG	GTTTTGTGGG	AAAGTTATAAG	TTCGGATCCG	5940
TTCTGACCGG	TCACCAAGTAC	CGATTGTCGA	CTTACGCTCA	GCCTCCTGCT	CATCTTGAGC	6000
AATCTGAGCA	AAGTGGTCAG	CAACACGGGC	ACGGATGATT	TTCATGGCCT	TCTCACGGTT	6060

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CTTCTGCTGG	GTACGTTCTT	CCTGCATCTC	AACCTTGATA	TTGGTTGGCA	AGTGAACGAT	6120
ACGAACGGCA	GTCGCAACCT	TATTGACGTT	CTGTCCACCA	GCACCAGAGG	CGTGATAGAT	6180
GTCGACACGA	AGGTCTTTG	GATCAATGTC	GTATTCAACC	TCTTCAACTT	CTGGCATAAC	6240
AAGAACTGTC	GCTGTCGAAG	TATGAACACG	GCCTTGCGTT	TCTGTCACAG	GAACACGTTG	6300
CACACGGTGG	GCACCTGATT	CATACTTAAG	CTTAGAGTAT	ACAGACTGAC	CTGAAACCAT	6360
AGCAACCACT	TCTTTAAAAC	CACCGACACC	ATTCATAGAG	GCTTCCATGA	CTTCAAAGCG	6420
CCAACCTTGG	GCTTCCGCAT	ACTTTTGGTA	CATAGTTAGC	AAATCTCCAG	CGAAAAGTGC	6480
CGCTTCGTCT	CCACCACTG	CTCCACGGAT	TTCAAGGATG	ATATTCTTGT	CATCGTTGG	6540
ATCCTTTGGA	AGGAGCAAA	TTTCAGTTT	TTCTTCATAT	TCTTCTTTT	CAGCCTTGGC	6600
ATCTTTGAGT	TCTTGCTTGG	CCAATTCTTC	CAAGTCCGCA	TCTCCGCCTG	ATTCCCTTAAT	6660
CATCTCTTCG	GCATCGACGA	TATTTGAAG	GACTTGTAA	TACTCACGGT	AGGCTATTAC	6720
GGTGTACCGA	TTGGAAGCTT	CTTCTTTGTA	AAGCTCCATA	AAACGTTGG	TGTCTGAAAC	6780
GACATCAGGG	TCACTCAGCA	ATTCTCCTAA	TTCTTCATAA	CGGTCTTCTA	CAACTTGTAG	6840
TTGATCATAG	ATGTTCATTT	TTTCTCCTTA	TTTCTCAATT	GTAAATCAT	AGATTGCTAC	6900
TACTTCATTC	TCGGATATTT	CCCCAGTTTC	TTTAAATCCA	AAACTGAGGT	AAACAAATCT	6960
TGCCCTGTTCA	TTTCTGGTT	CATAAGACAA	CCAAAGTTTA	TTGCTTAAAC	CTGCTGGCGC	7020
TGTTCGAAC	TAGTCTAGTA	CTTTATCCAT	AAATGGTTA	AAATATCCTT	GATTTGAAA	7080
ATTCTTATCA	ATCATAAAAC	AAAATAGTAA	ATAATTCCA	CTACTAATTC	CGATTTTTT	7140
ATCATAAGCT	ATCATCACAA	AACCTATAAT	TGCATCATTA	TCATAAACTG	CCAATGGAGC	7200
TACAAAATCT	CCATTTTAG	TGTAGACGTA	TGCTTCAGCT	AAACTAATTG	CGTTGGTTGC	7260
AATGAATTGT	TTTTGATATT	CCTTGACATC	CAAATTAAA	ACATCAAAT	AAATTTCCAT	7320
TGTAACATCT	CTTAGTTCAA	TTGTCTAGT	TTTGCTCCTT	GTTAGAGGTT	ATCATTGGCG	7380
CAAATAATG	TTTACGGCAA	ACTGAGATAT	AGGTTCGTT	ACCACCAATC	TGGATCTGTT	7440
CTCCATCGTA	AACGGGCAGT	CCATCCTGTG	TTCGCAACAC	CATGGTCGCC	TTTTCTTGC	7500
AATACTGACA	GATGGTCTTG	ATTTCGTCAA	TCTTGTCTGC	AAAAAGCAAG	AGATATTG	7560
AACCTTCGAA	CAATTTCATTG	CGAAAGTCAT	TTTTCAAGCC	AAAAGCCATG	ACGGGTATGT	7620
CTAACTCGTC	CACAAACACGA	GCTAGGTCGT	AAACATGGTG	GCGTTTGAGA	AACTGGGCTT	7680
CATCGACCAA	AACACACTAA	GGTTTTCTG	GTAGGTCTCG	GATATAGCCA	AAGATATCCG	7740
TTGTTTCCTC	AATCGCAAGG	GCAGGGCGTT	TCATGCCAAT	TCGACTCGAC	ACATAGCCAA	7800
CGCCGTCACG	CGTATCCAGA	GCCGAGGTCA	TAATCACAAC	ACCTTTCCCT	TGCTCCTCGT	7860

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AGTTATAGGC CACTTGAGA ATCTCAATCG TTTTACAGA GTTCATGGTC CCATAACGAT	7920
AGTACAAC TGCCATGTTT CTTGCTTCAC GTCCATTCTC AAATTTTGC TACATTCTAG	7980
TATATCATAA TTTCTTAAG CTTTAAACGG CAAAATGTGG TAAAATAGAA GAAATCAAAA	8040
ACTAGTGGAG GAAGCTATTA TGCCATTGT ACGCATCGAT TTATTTGAAG GACGCACGCT	8100
CGAGCAAAAG AAAGCTCTG CTAAGGAAGT AACGGAAGCA GTTGTCCGCA AACTGGAGC	8160
CCCTCAATCT GCTGTCCATG TCATCATCAA CGACATGCCA GAAGGAACCTT ACTTCCCACA	8220
AGGGAAATG CGTACTAAAT AAGCTAGCTT AAGCAGAATT GCTTAGGCCTT TTTCAATCTC	8280
CAAGTAGCAT TCATTGAAGA AATATCCTAA ATTTGTTACA ATTTGAAAAG AAACTTGGAG	8340
AATTCCAAG AAAAGAGCTA TTAATTAAG GAAACATTAT GATTACACGT GAATTGATA	8400
CCATCGCTGC TATCTCTACT CCACTAGGTG AAGGGCTAT TGGTATTGTC CGCCTGAGCG	8460
gAACAGACAG TTTGCTATT GCGAAAAGA TTTTAAAGG AAAAGACTTG AACAAAGTTG	8520
CCAGCCACAC TCTCAACTAC GGTCACATTA TTGATCCTCT GACTGGTAAA GTCATGGACG	8580
AGGTTATGGT TGGGCTATG AAGTCTCCAA AGACCTTCAC TCGTGAGGAT ATTATCGAGA	8640
TTAACACCCA CGGTGGGATT GCGGTGACCA ATGAAATTCT CCAGCTAGT ATTCTGTGAG	8700
GGGCTCGTTT GGCAGAACCT GGTGAATTAA CAAACACGTGC TTTTTAAAC GGTCGCGTAG	8760
ACTTGACACA GGCAGAGGCT GTGATGGATA TCATCCGTGC CAAGACTGAC AAGGCCATGA	8820
ACATGCGGT CAAACAATTA GACGGCTCCC TTTCTGACCT CATTAACAAT ACCGGTCAAG	8980
AAATCCTCAA TACACTTGCC CAAGTTGAGG TCAATATCGA CTATCCTGAG TATGACGATG	8940
TTGAGGAAGC CACTACTGCT GTTGTCCGAG AGAAGACAAT GGAGTTTGAG CAATTACTAA	9000
CCAAACTCCT TAGGACAGCA CGTCGTGGTA AAATCCTTCG TGAAGGAATT TCAACGGCTA	9060
TCATGGACG TCCCAACGTT GGGAAATCAA GCCTTCTCAA CAACCTCTTG CGTGAGGACA	9120
AGGCTATCGT AACAGATATC GCTGGGACAA CACGAGATGT CATCGAAGAG TACGTCAACA	9180
TCAATGGTGT ACCCTCTCAA TTGATTGATA CAGCCGGTAT TCGTGAACG GATGATATCG	9240
TTGAACAAAT TGGAGTTGAG CGTTCGAAAA AAGCTTTAA GGAAGCTGAC CTAGTTCTGC	9300
TAGTACTAA CGCTAGTGAA CCACTAACCG CCCAAGATCG CCAACTCCTA GAAATCACTC	9360
AGGAGACTAA TCGCATTATT CTTCTTAACA AAACTGACCT GCCTGAAACG ATTGAAACTT	9420
CGGAACCTACC TGAAGATGTC ATCCGCATT CAGTTCTTAA AAATCAAAC ATCGATAAAA	9480
TCGAAGAGAG AATCAACAAAC CTCTTCTTG AAAATGCTGG TTTGGTTGAG CAAGATGCTA	9540
CCTACTTGTC AACGCCCGT CACATTTCCCT TGATTGAGAA GGCGTTGAA AGCCTACAAG	9600

714

CTGTTAACCA AGGTCTTGA	CTAGGGATGC CAGTTGACTT GCTTCAGTT GACTTGACCC	9660
GTACTTGGGA AATTCTAGGA	GAAATCACTG GAGATGCTGC TCCAGATGAA CTCATCACCC	9720
AACTCTTAG CCAATTCTGT	TAGGAAAAT AAGAAAATC CATGATCCTT CATTGGTCA	9780
TGGATTTAG GTTCTATAAT	ATTTGTAGTG GGTAATCCA CTATAGATAT TATGGAGCCT	9840
ATTTTATTGT AGAAAAAAAG	TCCCATATGA CCTATAATGA AAAGCGACAA AACAACTCAT	9900
TAGAAAAGAT CATATGGAAC	AATTACATTT TATCACAAAA TTACTAGACA TTAAAGACCC	9960
TAATATCCAG ATTTTAGACA	TCATCAATAA GGATACACAC AAGGAAATCA TCGCCAAACT	10020
GGACTACGAC GCCCATCTT	GCCCTGAGTG CGGAAACCAA TTGAAGAAAT ATGACTTTCA	10080
AAAAACCTTC TAAAATTCTT	TATCTTGAAA CGACTGGTAT GCCCACTAGA ATTCTCCTTA	10140
GAAAGCGTCG ATMCAAGTGC	TATCACTGTT CAAAATGAT GGTCGCTGAA ACTTCTATCG	10200
TCAAGAAGAA TCACCAAATC	CCTCGTATCA TCAACCAAAA GATTGCTCAA AAGTTAATTG	10260
AAAAGATTTTC TATGACTGTAT	ATPGCCCATC AGCTTCCAT CTCAACTTCA ACTGTTATTC	10320
GTAAGCTCAA TGACTTTCAC	TTTAAACATG ATTTTCTTG TCTTCCTGAG ATTATGTCTT	10380
GGGATGAGTA TGCTTTACA	AAAGGGAAGA T	10411

(2) INFORMATION FOR SEQ ID NO: 90:

- (i) SEQUENCE CHARACTERISTICS:
- (A) LENGTH: 2393 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 90:

GTTTGGGTT CTGAAATTA	TCAGATGGTT GGAAAAGCCG TCCACATCAA GATAGTGTTC	60
GGAGATTAA GTTAAATTG	AAGAAACTAA CACAGAGGAA ATGGAGTATA GACCTAACAA	120
GACGTATTGA GCAACTGAAT	TTGTCTATTG GAGGATGGAT AACTATTGC TCATTGGAA	180
ATATGAAAAG TATAGTCGCC	AGCATAGATG AGCGCTTGCG TACTCGCTA CGAGTGATTA	240
TCTGGAAGCA ATGGAAGAAG	AAATCGAGAC GATTATGGGG ATTGCTTAAG TTAGGACTTC	300
CTAAATGGAT AGCAGATAAG	GTATCTGGCT GGGGCGACCA TTATCAATTA GTAGCTCAGA	360
AGTCGGTACT TAAACGTGCT	ATATCAAAAC CAGTCCTGGA AAAACGTGGA CTGGTTTCGT	420
GTGGGATTA TTACCTTGAA	CGACATGCGT TAAAAGTTAG TTGAACCGCC GTATGCCAA	480
CGGCACGTAC GGTGGTGTGA	GAGGGGCTAG AGATTATCCC CTACTCGATT AACTCCCCTG	540
AAATTATTT TAATTATGCA	AATTTCACGT ATTTTGATG CTGAGACGAC GATCCTGGGA	600

715

ACTTTTCAGA TATTTTTTG ACTATCTAA TCTATCATTA GAAAAGCTTA GAGGCCAAA	660
GGATTTGAGC GTTTTCTGA TTTTTAAGAC TTTTTCCAGT CTCTTTCTG ATTGAAGATG	720
TAATTATTCT ACTAACTAAC TAACCTCTTA GTACTAGCCA ACAACGATAA TCATAATTCC	780
TCCTAAAATT AGGAATAATA AAGGCAATAG TTTTGTTT TTCACTGTAA AACACCTCACT	840
TTTGTCTCT GCTATTTAT GCTAAAATAT TAAAAATCAA ATTTAATTCC AAAGTTGTA	900
ACTAAAGGG GAGCGCTACA TGTCTAAC CTTGTCAAG TTGTTAGTCT CTCAATTATT	960
TGCAAATTAA GCAGATATT TCTTTAGAGT AACAACTATT GCTAACATAT ACATTATTTC	1020
AAAATCAGTA ATTGCCACAT CACTAGTTCC TATCTTAATA GGAATATCCT CTTTGTTGC	1080
GAGCTTTTA GTTCCGTTGG TTACTAAAAG GTTAGCGCTA AATAGGGTT TATCTTATC	1140
TCAATTGGA AAGACTATAT TATTGGCGAT ACTGGTAGGA ATGTTACCG TAATGCAATC	1200
CGTAGCGCCT TTGGTGACCT ATCTATTTGT TGTTGCAATT TCCAACTAG ATGGTTTG	1260
AGCACCCGTT TCCTATGCTA TTGTGCCACG CTATCGACCC GATTGGGTA AGGCTAATT	1320
AGCCTTATCA ATGACTGGTG AAGCTGTTCA ATTGATAGGT TGGGGATTAG GTGGACTCTT	1380
GTTGCAACA ATTGGTCTGT TACCTACCAC GTGTATCAAT TTAGTCTTGT ATATCATTTC	1440
TAGCTTCTG ATGTTATTTT TTCCTAACGC TGAAGTGGAG GTGTTAGAGT CAGAAACTAA	1500
TCTTGAAATT TTGCTCAAAG GTTGGAAAGTT AGTTGCTAGA AATCCTAGAT TAAGACTTT	1560
TGTATCAGCA AATTATTGAA AATTTTTTC AAATACGATT TGGGTTCTT CCATTATACT	1620
TGTTTTGTA ACGGAGTTAT TAAATAAAAC GGAAAGTAC TGGGGATATT CTAATACAGC	1680
ATACTCTATT GGTTTATAA TTAGTGGCTT AATTGCTTT AGGCTATCTG AAAAGTCCT	1740
TGCTGCTAA TGGGAACCCC AATTATTACAC CCCAAATCTA AAAACCATCC AGAACCTTG	1800
CCTTAGCTTA GATCCTGGAT GGGTTCTTTT TTCACCCAAAT GGGTGTCTT TACTAGACAA	1860
AAAAGAGTTT CCCCTTTATG GTATAAGTGT AGAAAAAAAC ACAAAAAGAA AGGAAACTCA	1920
CATGAACAGT TTACCAAATC ATCACTTCCA AAACAAGTCT TTTTACCAAC TATCTTCGA	1980
TGGAGGTCA TTAACCCAGT ATGGTGGCT TATCTTTT CAGGAACCTT TTTCCAGTT	2040
GAAACTAAAA GAGCGGATT CTAAGTATTT AGTAACGAAT GACCAACGCC GCTACTGTCG	2100
TTATTCGGAT TCAGATATCC TTGTCCAGTT CCTCTTTCAA CTGTTAACAG GTTATGGAAC	2160
GGACTATGCT TGAAAGAAT TGTCAGCTGA TGCCTACTTT CCAAAATTGT TGGAGGAGG	2220
GCAGCTTGCT TCACAGCCAA CCTTATCCCG TTTTCTTCC AGAACTGACG AGGAAACAGT	2280
CCATAGTTG CGATGCCCTCA ACCTTGAATT GGTCGAATT TTTTACAGT TTCACCAGCT	2340

716 AAACCAACTC ATTGTAGATA ACGATTCTAC CCATTTACACA ACTTATGGCA AGC 2393

(2) INFORMATION FOR SEQ ID NO: 91:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 4762 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: double
(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 91:

717

TGGTTTTAAG ATTGGAAATG CAGAGGTAGA GAGTCCCAAG TCTATCCAGA CTGCGACAGC	1440
ACAGATTCT CAAATCATTC CCAACGTTGC TTCTAGCCAG TACGGTGGCT GTTCAGCTGA	1500
CCGTATCGAT GAAATTTGG CGCCTTATGC AGAGAAGAAT TATCAAAAAC ATCTCAAAGA	1560
TGCAGAAGAG TGGGTATTGC CTGAAAAACA GGAAGATTAC GCTTGGAAAGA AAGCGCAAAA	1620
GGACATCTAC GATGCCATGC AATCTCTTGA GTATGAAATC AATACTCTCT TCACTTCAAA	1680
TGGACAAACA CCTTTTACTT CGTTAGGTTT TGGTCTGGGA ACCAGTCGTT TTGAACGAGA	1740
AATTCAAAAA GCTATTTAA ACATTCGCAT CAAGGGTCTT GGTCAGAAC ACCGTACGGC	1800
TATCTTCCTT AAACCTTATCT TTACGCTTAA AAGAGGCCTC AACTTAGAGG AAGGAACCTCC	1860
CAACTATGAC ATCAAGCAGT TGGCTCTAGA GTGTGCAACC AAGCGGATGT ATCCAGACGT	1920
CTTGTCTTAT GATAAGATTG TTGATTGAC AGGTTCTTTC AAGGTGCCTA TGGGCTGCCG	1980
TTCTTTCTT CAAGGGTGGAGGATGAAAA TGGTGTAGAA GTCAATTCAAG GTCGCATGAA	2040
TCTGGGTGTT GTGACGGTTA ATCTGCCTCG TATTGCTCTT GAGTCTGAAG GTGATATGAA	2100
TAAGTTCTGG GAAATCTTCA ACGAGCGAAT GAATATCGCA GAAGATGCTC TTGTTTACCG	2160
TGTCGAACGC ACTAAAGAGG CGACACCAGC GAATGCTCCT ATTCTTTATC AGTACGGTGC	2220
TTTTGGCCAT CGTCTAGGTA AAGAAGAAAG TGTTGACCAAG CTCTTTAAGA ATCGTCGTGC	2280
GACCCTTTCG CTGGGCTATA TCGGCTTGTA TGAAGTAGCG ACAGTTTCTT TTGGTAACAG	2340
CTGGGAAAGT AATCCAGATG CTAAGGAATT CACGCTAGAC ATCATTCACG ATATCAAACG	2400
CCGTGTAGAA GAGTGGTCAG ACCAATATGG CTACCATTTTC TCTATCTACT CAACACCATC	2460
CGAAAGTCTG ACAGACCGTT TCTGCCACT AGATATAGAC AAGTTGGCT CTATTCTGA	2520
TATCACAGAC AAGGAATACT ACACCAACTC TTTCCACTAC GATGTTGCTA AAAATCCAAC	2580
ACCGTTGAA AAATTGGACT TTGAGAAAGT CTATCCGGAA GCAGGTGGT CAGGTGGTT	2640
CATCCATTAT TGTGAGTATC CAGTCCTTCA GCAAATCCA AAGGCCCTGG AAGCTGCTG	2700
GGATTATGCT TATGACCGTG TAGGCTATCT AGGCACCAAT ACTCCGATTG ACCGTTGCTA	2760
CAAGTGTGAC TTTGAAGGGG ATTTGAAACC AACTGAGAGA GGGTTGCTT GTCCAAACTG	2820
TGGCAATAGC GACCCCTAAAA CAGTAGATGT GGTGAAACGA ACTTGTGGCT ACCTAGGTAA	2880
TCCTCAAGCA AGACCGATGG TCAACGGCG TCACAAGGA ATCGCTGCCG GTGTCAAACAA	2940
TATGAATGGT TCAACGATTA AAATAGCTGG GCATCAAGTA ACAAAATTAGA AAGAAATGAA	3000
ATGGGAAAT ATCAACTAGA CGATAAGGGG CGCGCACAAG TGACCCGTTA TCACGAGAAA	3060
CACTCTAAAG GTGGAGCTGG TAAGAAAGAA CGCTTGCTTA GCTTCAGAGA ACAATTCTTA	3120

718	
AAACAAGAAC A GAAAAAAATA AAAGTGAGAG CCAGCTCTCG CTTTCTCAT AGTGGGAGGT	3180
AAGGATGGAA TTACGCAGAC CAAGATTAGC GGATAAAGAAA GCTGTTTAG ATATGATGAC	3240
AGAGTTGAA AAATTCAGT CGCCTCACGA CGGCGGTTTC TGGGATACAG AGAACTTTGT	3300
GTATGAAGAC TGGTTAGAAA GCAATCAGGA ACAGGAAATG GGGATTAATC TGCCTGAAGG	3360
ATGGGTTCT GCAATTCACT TAGTGGCTTT TTCTGAGAAA GGTCAAGCAG TTGGATTCT	3420
TAATCTCCGG TTGCCCTCA GTAACTTTCT ACTAGAAGAA GGTGGCCACA TTGGCTACTC	3480
CATTCGTCCA TCTGAAAGAG GCAAGGGTTA TGCAAAAGAG ACTCTCCGTC AGGGCTTGCA	3540
AGTTGCTAAG GAAAAGAAC A TCAAGAAAGC TCTGGTGACC TGTAGTGTGA ATAATCCTGC	3600
TAGCAGAGCA GTCATTCTAG CAAATGGTGG AATATTTGAG GATGCTCGCA ATGGAGTCGA	3660
CGCTTATTGG ATAGAGGTAG CGAATGAATA ATCCAAAACC ACAAGAATGG AAAAGCGAGG	3720
AACTTAGTCA AGGTCTGTATC ATTGACTACA AGGCCTTTAA CTTTGTGGAC GGCGAAGGCG	3780
TGCGCAACTC TCTCTATGTA TCAGGCTGTA TGTTTCACTG CGAGGGATGT TATAATGTTG	3840
CGACTTGGTC TTTTAATGCT GGCATTCCCT ATACAGCAGA ATTAGAAGAG CAGATTATGG	3900
CAGACCTTGC CCAACCCSTAT GTTCAAGGCT TGACTTTGCT GGGAGGGAG CCTTTCTCA	3960
ATACTGGGAT TCTCTTGCCA CTTGTTAACG GGATTCGGAA GGAATTGCCA GACAAGGACA	4020
TCTGGTCTG GACCGGCTAC ACTTGGGAAG AAATGATGTT GGAAACTCCA GATAAACTGG	4080
AATTCCTTGTC ACTGATTGAC ATTCTTGTCG ATGGAAGATA TGATCGAACT AAGAGAAATC	4140
TTATGCTCCA GTTTCGAGGT TCATCTAACCC AACGAATTAT CGATGTGCAA AAATCGCTCA	4200
AAAGTGGGCA AGTAGTGATT TGGGACAAGC TCAATGACGG AAAAGAAAGC TATGAACAGG	4260
TGAAGAGAGA ATGAAAGAAA AGGACTTAGT AGACCAACTA GTCTCAGAGA TCGAGACGGG	4320
GAAAGTCAGG ACACTGGGAA TATACGGTCA TGGAGCTTCA GGTAAATCAA CCTTTGCACA	4380
GGAAATTGTAC CAAGCTTTAG ATTCTACTAC AGTAAATTG CTAGAGACAG ATCCTTATAT	4440
CACCTCAGGA CGCCATCTGG TAGTACCCAA GGACGCCCG AATCAAAAGG TGACAGCCAG	4500
TCTGCCAGTG GCCCATGAAC TGGAGAGTTT GCAGAGAGAT ATCCTTGCTT GCAGGGGGT	4560
ATGGATGTCT TGACAATTGA AGAACCTTGG AAGGCTAGTG AGGTCTTGTC TGGAGCCAAA	4620
CCAATTTGTA TTGTCGAAGG GATGTCTGTT GGCTTCTAC CCAAGGAACT CTTTGAAAAA	4680
ACCATCTGTT TCTACACGGA TGAGGAGACC GAATTAAAGC GACGCCCTGC TAGAGATACG	4740
ACTGTGAGAA ATCGCGATGC GG	4762

(2) INFORMATION FOR SEQ ID NO: 92:

(i) SEQUENCE CHARACTERISTICS:

719

- (A) LENGTH: 3832 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 92:

GATGCAGGTT TCGACCCACA TATTCCAGAA AATTACTTTA AAGATGATGA TGTTAACAG	60
GTACCTGTC TTTGTTGGTC TTCATCTGCA GCCCTCTTT TCAGTAATTG GGTAGACCAT	120
GGCGTCTATC AGGAGACGCC TTTGATTGG AGAAAGATAG AAGATGATGC ATCTGCATAT	180
GGGTATTTAT AAGAGGAATT ATGACATATT TAGACGCTTT TAAATCAGGT ACCTTGGTTT	240
TACCGAGTGC CCTGCTCTTG CATTAAAGG AACTCTTCC TTCTAGCGAC GATTTCTGG	300
TTTGGCAATT TTTCTATTTG CAAAATACGA CAGGCTTAGA AGAAATGTCG CCAAGCCAGA	360
TTGCTGAAAG GATTGGCAAG GAAATTTCGG ATGTCAACCA GTCCATTCT AATCTGACGG	420
AAAGGGGACT GCTCCAGTAT CGTACTATCG AATTAAATGG CGAAATTGAA TTGCTCTTG	480
ATGCTAGTTT GGCCCTGGAA CGTTGGATG ACCTGTTGG AGCAGTTCAT TCAAGTTCA	540
ACCAAGCTAAC ACCTCAAAAC CAGCTCAAGG ATTGTTGGAA AACCTTCCAG CAGGAGTTGG	600
GACGATTGTT GACGCCTTTT GAGATTGAGG ATTTGACCAA GACACTAAAG GAAGATGGAA	660
CCAGTGCTGA CTTGATTAAG GAGGCTCTTC GTGAAGCTGT TTTGAATGGAA AAACCAAAC	720
GGAAGTACAT TCAGGCGATT TTGAGAAACT GGCGCCATGA AGGAATCAAG AGTCTCCCTC	780
AAATTGAGGC CAAGAGAGCA GAAAGAGAAG CAAGCAATCC TCAGTTGACA CAGGTATCTG	840
CAGATTTCAT AAATGCCATG GATCTCTGGA AGGATTAATC CATGCAAGTA GGCTTGAAAT	900
CCGAGTAAGA TTTGCAAGCT GTGTATAATT GTGATAGAAAT AAATAGAAAA TAAATTGAAA	960
AAAGAGGTAT GTGAAATGTC ACGTAAACCA TTTATCGCTG GTAACGGAA AATGAACAAA	1020
AATCCAGAAG AAGCTAAAGC ATTGTTGAA GCAGTTGCAT CAAAACCTCC TTCATCAGAT	1080
CTTGGTTGAAG CAGGTATCGC TGCTCCAGCT CTTGATTTGA CAACTGTTCT TGCTGTTGCA	1140
AAAGGCTCAA ACCTTAAAGT TGCTGCTCAA AACTGCTACT TTGAAAATGC AGGTGCTTTC	1200
ACTGGTAAA CTAGCCCACA AGTTTGAAA GAAATCGGTA CTGACTACGT TGTTATCGGT	1260
CACTCAGAAC GCCGTGACTA CTTCCATGAA ACTGATGAAG ATATCAACAA AAAAGCAAAA	1320
GCAATCTTG CGAACGGTAT GCTTCCAATC ATCTGTTGTG GTGAATCACT TGAAACTTAC	1380
GAAGCTGGTA AAGCTGCTGA ATTGCTAGGT GCTCAAGTAT CTGCTGCATT GGCTGGATTG	1440
ACTGCTGAAC AAGTTGCTGC CTCAGTTATC GCTTATGAGC CAATCTGGGC TATCGGTACT	1500

720	GGTAAATCAG CTTCACAAAGA CGATGCACAA AAAATGTGTA AAGTTGTTCG TGACGTTGTA	1560
	GCTGCTGACT TTGGTCAAGA AGTCGCAGAC AAAGTTCGTG TTCATAACGG TGGTCTGTT	1620
	AAACCTGAAA ATGTTGCTTC ATACATGGCT TGCCCAGACG TTGACGGTGC CCTTGTAGGT	1680
	GGTGCAC TTGAAGCTGA AAGCTTCTTG GCTTGCTTG ACTTTGTTAA ATAATCAGTA	1740
	AGTAGCAAAA GCTAGGTGGA ACAGCATTCA GATGTCCTGT ACATTTTTA TAGGAGAGAA	1800
	AGATTGAAAA CAAAAATTGG ATTAGCAAGT ATCTGTTTAC TAGGCTTGGC AACTAGTCAT	1860
	GTCGCTGCAA ATGAAACTGA AGTAGCAAAA ACTTCGCAGG ATACAACGAC AGCTTCAAGT	1920
	AGTTTCAGAGC AAAATCAGTC TTCTAATAAA ACGCACAGCA CGCGCAGAAGT ACAGACTAAT	1980
	GCTGCTGCC ACTGGGATGG GGATTATTAT GTAAAGGATG ATGGTTCTAA AGCTCAAAGT	2040
	GAATGGATTT TTGACAACTA CTATAAGGCT TGTTTTATA TTAATTCAAGA TGGTCGTTAC	2100
	TCGCAGAACATG AATGGCATGG AAATTACTAC CTGAAATCAG GTGGATATAT GGCCAAAAC	2160
	GAGTGGATCT ATGACAGTAA TTACAAGAGT TGTTTTATC TCAAGTCAGA TGGGGCTTAT	2220
	GCTCATCAAG AATGGCAATT GATTGGAAAT AAGTGGTACT ACTTCAGAA GTGGGGTTAC	2280
	ATGGCTAAA GCCAATGGCA AGGAAGTTAT TTCTTGAATG GTCAAGGAGC TATGATGCAA	2340
	AATGAATGGC TCTATGATCC AGCCTATTCT GCTTATTTTT ATCTAAAATC CGATGGAAC	2400
	TATGCTAACCC AAGAGTGGCA AAAAGTGGGC GGCAAATGGT ACTATTCAGA GAAGTGGG	2460
	TATATGGCTC GGAATGAGTG GCAAGGCAAC TACTATTTGA CTGGAAGTGG TGCCATGGCG	2520
	ACTGACGAAG TGATTATGGA TGGTACTCGC TATATCTTG CGGCCTCTGG TGAGCTCAA	2580
	GAAAAAAAAG ATTTGAATGT CGGCTGGTT CACAGAGATG GTAAGCGCTA TTTCTTTAAT	2640
	AATAGAGAAG ACAAGTGGG ACCGAACAT GCTAAGAAAG TCATTGATAT TAGTGAGCAC	2700
	AATGGTCGTA TCAATGATTG GAAAAGGTT ATTGATGAGA ACGAAGTGGA TGGTGTCAATT	2760
	GTTCGTCTAG GTTATAGCGG TAAAGAAGAC AAGGAATTGG CGCATAACAT TAAGGAGTTA	2820
	AACCGTCTGG GAATTCCCTTA TGGTGTCTAT CTCTATACCT ATGCTGAAAA TGAGACCCAT	2880
	GCTGAGAGTG ACGCTAACACA GACCATTGAA CTTATAAAGA AATACAATAT GAACCTGTCT	2940
	TACCCCTATCT ATTATGATGT TGAGAATTGG GAATATGTAA ATAAGAGCAA GAGAGCTCCA	3000
	AGTGATACAG GCACTTGGGT TAAAATCATC AACAAAGTACA TGGCACACGAT GAACCAGGCG	3060
	GGTTATCAAA ATGTGTATGT CTATAGCTAT CGTAGTTTAT TACAGACGGC TTTAAAACAC	3120
	CCAGATATTT TAAAACATGT AACTGGGTA CGGGCCTATA CGAATGCTTT AGAATGGGAA	3180
	AACCCCTCATT ATTCAAGAAA AAAAGGTTGG CAATATACCT CTTCTGAATA CATGAAAGGA	3240
	ATCCAAGGGC GCGTAGATGT CAGCGTTTGG TATTAAGCGA TGATTTGAAA GAGGGATGTG	3300

721

ATAGTAGCAC CCTCTTTTC TTTGTTTAT GATAGTCAT CCTCGAGTAA ATTCAAGTTC	3360
TTGCTCGAA ATGAAGCTTA TATAGTAGAT TGAATATAGA CAAATACCTT GTGATTGGTA	3420
AAACATTTA GAAATTCAATTACCTT TACCTTTCTT AATCGACTTG GTTTCATCTT ATTTCAATCT	3480
ATTATAGTAT TGGGGAATTTC TTCAAAACCA CATCAGCTTG GTCAGTCTA CCTGCGACCT	3540
CAAAACTGT GCTTGGTCA AGCTGGTTT AGTTTCCTAG TTTGCTGATG GATTTCCATT	3600
GACTATAAGC ATCCAACCCT CTTTTGTCT TCTAAAGAAT TCTTAAATTCA TCAGTCTATT	3660
GCAACTTTTC TCATATAAGT TCTTTGTCTT GCTATTGGTT TTCCTTAGTA GTATACTAAG	3720
GTAGTAATCA TTAAGAAGTG GTTACAAAAA ATAATGAATG AGGTAAGAAA AATGGTAGAA	3780
TTGAAAAAAAG AAGCAGTAAA AGACGTAACA TCATTGACAA AAGCAGGCC GG	3832

(2) INFORMATION FOR SEQ ID NO: 93:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 10690 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 93:

TGAAAAAAATC CTCATGAACC TGGCGCCAAT AGACAAGTGT CTTGTTTCCC TCACCTTCCT	60
TATAGGCATG GTCAGCTGAC ACTCGATTGA AGGGTTAAC AGAAACCTTT GTAATTCCAA	120
CAATGCAGAC AGCCTGATTT TGACTATCTA AAATGACATC GAAGGTCCT ACCTGGGGAA	180
GTGGTTGTC TTCTAGCACA TAGAGGTAT AGGCTGATCC TGTTGCTGTC TTTTCTCCCTT	240
TAAACACCAA ATCCGCTAAA AGGTCTGGTT CAACTCCAAA AGCCAGGCA TCGATTTCAT	300
CTCCGATCAA AGGATTGATT TGCTTGATT TATTCCACAT TTCTTGCGGT ATCATGGGT	360
CTCCTTGTA ATTTTTACT TTCTTCTTTT ATGTGTTAAC GATGATCTGG ATGGTCAATC	420
TCTAAATCAA AAATCTCTGG AATAGAACTG TAGTGGATAA TGCACTTGAT ACCCAACTGA	480
TTCATTTTT GTATGAAAGA AGTATTCAAGA TAGCCTGCTA CAGCAAATC AATCTTGTTC	540
TTTCTTGCTT TATCCTGCAT ATCTCTTAGC ATATCTAACAA TTATTGGACT TTCCATATCA	600
TGCCATTGAC TGTCTCTCAT AGTCGCAAAA ACAAGGAAG TCAAATCATT CATTCCAAC	660
ACAATTTTG AAATGCCGT TTCCAGTATA CTAGATAAGT CAAATACCC TGACGGTAAT	720
TCAATCATCG TTCCGACTTT CCCAGTAAA CCCTGCTGAC GCAATACTGT AATAGCTTGT	780
TTTAATTGGT CGGCATCATT GACAAAAGGA AAGATAACAG ATAGATTGGG GTTGGTTGA	840

722

TAAACTTCTG	TAACGACATG	TGCTTCAGCC	TGAAATTCA	CCAAACACGC	CAGTAAACGC	900
CTAGTTCTC	TATAGCCAA	CAAGGGATGC	CCTTCGTCAA	AAAACTTTT	AGTCCCCACT	960
AAACAATTGG	CTTCTGTATT	CGTTAATTCA	GTAAAACGAT	ACCAAACCTC	CTTACCTAAG	1020
TAAAAGGAGC	AAATAGTATC	AAGATAATCT	TTCACAAATT	CCTGACAACT	TTGTAATAGT	1080
ATATTTGAT	TGAGCTCTCT	CAATAAGTAT	TCCCCACGAA	TCATGCCGAC	GTGGTGAAAT	1140
AGTTGAGGAT	AAATTTTTTC	AAGAATTTTT	TCGCCACTAA	GGGCAAGTTG	ATTTCTCATC	1200
ATTCACCTTC	CAATTCATGT	AAGAAGTCTT	GTCCAGTTCT	GGAAATCCTA	ATAATTCTAGA	1260
CTTAACCTTC	AAGACTAATG	GCGATGCATT	TTCTTCTGTA	ATCTCTTGA	TATCCATCCA	1320
AATATATCCA	AGTGAATCAT	TCGCACCATC	AGACACAGCT	TCCGAATCG	TAACATTGAGG	1380
TGCACTCTCA	TTCATAACTT	CATCATACAA	GGCTATGACA	TGGTGAACCA	AAAAATTTTT	1440
TAACTCTTCC	CTGACGAAAA	CATCGTAGAT	TCGAGGATT	GAGTAGCTTC	TAACAGTAAA	1500
TCCCGTCTCT	TCCATAACTT	CTCTAGTCAG	CGTTTCCGTC	AGTCCTTCAC	CAAGTTGCTG	1560
ACTGCCTCCA	GGTAGATCAT	ACCGATGTTG	ATAAGGGCCT	CTCGTTTTT	CAATGCAAAG	1620
TAACATTCCA	TTTTCAAAGC	AAACACAGTA	GACCCCAAAG	TGATTTTGA	TTTCCATCCA	1680
ACTCCTCCTA	CTTCAAAGAC	CAGCCACCAC	CTATTGTCAA	GATTTGTCCT	TGCATGGCGC	1740
TGCGTTTCC	ACTTGCTAAA	AAAAGACTAA	GCTCTGCTAT	TTCCCTGCGC	TCAATCCAGC	1800
GCTTGATTGG	GGTTTCACTA	GCCACCCAGT	CAGCCAAACC	ACCTGGTCA	AAATCCGCAG	1860
CGGTCACTAGC	TGTCTTGACT	GCTCCTGGAG	CGATACAAA	GACCTGAATC	CCAGCTTCAG	1920
CATAGTCTAG	AGCCAACACTG	TTGGTGAAGC	CAGCCAAGGC	ATGCTTGGAT	GAAGTATAGG	1980
CGTGACCACC	TCCACCTGCT	AGGCTAGAAC	CAATGGAACA	CATATTGATG	ATGATTCCT	2040
TTTTATTTTC	CAGCATTGT	GTCAAATAAT	ACCGAGTCAA	CTCTACTGGA	ATAATGTAGT	2100
TGATTTCAAA	AACTCTTGA	ATGTCCTGCG	CCGTTGTTTC	CAACAGTGGT	TTGTAATCAT	2160
CCAAAACCTCC	AGCAGTATTA	CACAAACAT	CCACCTGAGG	GCACCAGTCA	AAAATAGTT	2220
CCAAGTCCAA	GGTCAAATCT	CTCTGTAAAA	AGCGAAAATC	ACCCCTCTAAG	AGTGGCTTT	2280
CACCTGGTC	AACTCCATAA	ACTTGATAGC	CCTCTCTAA	AAAGAGGCCA	GCTTGAGCCA	2340
ATCCGATCCC	TGAACACTACT	CCTGTAATGA	GTACACGTTT	AGTCATGCAC	TTCTACCCAA	2400
TCCGTTGCCA	AAACATCACA	AACTGTCGGG	CTCCACATGG	AAAAACCTTC	TCCTTCGCCA	2460
GAAACGTTGA	TTAGGAAATA	AGGTGTCATT	TCAAGTGCAA	GCCCCATTTTG	CTCGATGGTA	2520
TCAAAGAGTT	GGACATAGTT	TTCCGCACCT	CCCCAACAG	TTCGTACATA	TTTTCTCTTA	2580
GCCTTTAAC	CAGGCAGGAT	CTCTTCAAAT	GTCATGTTT	TCTCCCTTAA	TTCTACATTC	2640

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TTCATTTAAT TATAGCAAAA AACCGCTTTA TACGGCTTTT TGAATGTGAG TTATTCAAAC	2700
CTGCTACTAC TTACGGCAAA TTATTCCCTG CAGCAAGATA AATTCATAC CATTCTTTC	2760
TTGTTAACGCT AAAGTTGCC GCTCGGCTAA CTCTCTCAA GTGCTTAGGA TTTGTTGTAC	2820
CTACGACTGC CTGCATTTT GCTGGATAAC GCAATATCCA AGAAATGGCA ATAGTTGAAG	2880
AGGTTACTCC ATATTTAATA GCTAAACGAT CAAGTACTTG ATTTAAAGCT TGAAATTCT	2940
CATTCCAAC AAAATTCCCT TAAAAAATACC CGAATTGTA GACAGACCAT GCTTGAATGA	3000
CCACATCGTG TAATTGGCAA TATTCAAAAA TGCTGCCATC TCGCATAGCT GCTTGACTAT	3060
CTTCCATATT AACATGAAAA GCTGATTCAA ATCCTGGAGT AAAAGCCGA CTCATTGTA	3120
GCTGATTAAC AGCTAACGGC TGCTTGACAT CTTTTTAAG CAACTCCATC ATCATAGGAT	3180
TTTGATTAGA AACTCCAAA TCTCGAACCTT TACCTTGTTT ATAAAGGAGA TTAAAGGCTT	3240
CTGCTACTTG GTCAGATTCC ATCAAAGCAT CTGGTCGATG AAGGAGCAAG CTATCTAGAT	3300
GATCAATCTT CAATCTTGC AAAATACCGT CTACTGATT TATAATATAG TCCTTAGAAA	3360
AATCAAATA GGTAATTCT TCAATGCGAA TGCCACATT GGACTGAATC CACATTTT	3420
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CATAAAATAC AGCCAAGTCG AAGGCATTGA TTCCAACAGA AAGTGCTGTT TCTACAAGCT	3540
CTTCAACTTC TTTTACAGAT TTATCTTTA TTCTCATCAT TCCGAGAACAA ATTCTGATA	3600
ATTCTTGTC ATCTTGACCA AGAGTTATGT ATCTCATCAA ATTTTCTCC TTTAATTCT	3660
AACATTCTTC CCTTCATTAT AACAAAAAAC CGCTTTGCAA CGACTTTTG ACTATACTTC	3720
ACTCCATTAA ATCTTCTTAA ACCCACGGAA CAAGACAAAG ATTCAAATAA AGAGGACAGC	3780
TAAAGGAATA ACTTTTGAA GGAAAACATT TGAAATTCCC ATCCACTCAT AATAACGGAG	3840
CAGAGAACCC ACCACAAGAT GGGCAATAAT CATACTGACA AATGGACGAA AGACCGCTTC	3900
TTTCCAATTG CAAATACCGA TAACTAGCGA AATCGTAAAG ACAGACAAAC TATCCAGGG	3960
AGCCGGAATA TAAAAGGCTC CTTCTTGAT GAAGCTTGCC ATTCCCTACAT ATCCTAAAC	4020
AACTAGAAGA ACTATAGTCC CAACAACAAT GTAAGTGCCA ATTTCTATT TAGGAGAATC	4080
TTGGACTAAA CTTCTTCGTA AAATTGTGGC CACAAGTCCA AATCCAATCA GAAAATAAG	4140
AAGTGCCTT AAAATGTGA GCAAATTGAC TGTAAAGAGA GGACCTTTAG AAAATCACT	4200
TAGTAGTTGA TAATAACGTA ATACCGCCAG GACAAGAATT GCGCTCAAAA GGGACTCTT	4260
GATAGAACTG CGAGGTGCTC CCTTGAGAAT CTCTTTCATT ATTTTTTAG GATTCTTAC	4320
TAGATAATCC TCTGCACTCA TGCCATCTCG TTCTGCTTCT GAGAAATCTA GCATCATCAA	4380

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ATAGATCTGC TCTCTGAGAT AGTCTTCATC ATAGAGAAAT CCAGCAAGAT TAAAACMTTC	4440
CCACAACTCC TCAAATACT TTTGATTCTC CTCAGAAAAC TCATGTAGCA AAGCGCTTGT	4500
TTCTTCGTAA TACTTCATTT TCTTCATGGT TTAACCCCCA TTCTTAATCC CTTCTACTTT	4560
TTGACTCAAA TCGTCCCATT GTTGCCAAA GACTGAGACA CGCTCTTCTC CTTCTTTCAT	4620
TAATGAAAAA TACTTCCGAT CTGGACCATC TGCGCACGGG CGCATGTCGC CTCTTATCCA	4680
TGATTTTTT TCTAACTTTT GCAACAAAGG ATAAATAGTT CCTGGAACGA TAGTATCAA	4740
TCCAGCCTCT CGCAAAGTCT GAACCAACTC ATAACCATAC CGCTCTTTT GACCAATCAT	4800
ATCCAAGACA CAACCTTCAA GAACACCTTT TAATAGCTGA GTTCTTCA TCACTTCTCC	4860
CTTCTAATCT ATTTGTAA ACCTACTAGT GACTTCACCT ATAGTATATC ACTTCTACAC	4920
TAGTTGTAA AGCATAATAG TTAATACTCT TCGAAAATCT CTTCAACCCA CGTCAGCGTC	4980
GCCCTACCGT ATGTATGGTT ACTGACTTCG TCAGTTTCAT CTACAACCTC AAAAACATGT	5040
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TCAGTTTCAT CTACAACCTC AAAACAGTGT TTTGAGCTGA CTTCGTCAGT TTCATCTACA	5160
ACCTCAAAA CATGTTTGA GCTGACTTCG TCAGTTTCGT CTACAACCTC AAAACAGTGT	5220
TTTGAGCAAC CTGCGGCTAG CTTCTTAGTT TGCTCTTGA TTTTCATTGA GTATAAATAA	5280
AAAACAGAA CTAGCCTGAA CTAGCCTGT CTACTTTTAC CCAATCACAC TTCCATTGG	5340
TACAGCTGGA TCAACTGTGA GAAGGGTTAA TTGCCCCATCA TGTTCACTG AGAGAACAT	5400
ACCCCTGGCTG ACATATTTT TCATCATTTC ACGTGGTTTG AGGTTACCAA CGATTTGAAC	5460
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ACGATCTCT CCATCACCAAGCG GAATTGAAGC AACTTATCTG AACCTTCTAC	5580
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GATTTCATCC TTGTTTAGTT TGAGCTCAAC TTCTGCCGA TTCCATTCTT TTTCGACTGC	5700
TGGTTTATTG CCTTCATTT GTTCCTTGAT ATAGGGAT TCTTCCTCCA TATTTAGACG	5760
TGGAAAGATA GGTGTTCTT TGGCAACTAC AGTCACATCT GCTGGGAAGT CAGCCAAACT	5820
CAAGTTTCA AGACTAGAAA CTTCTTCCAA ACCAAGTGA GTCAAAACTG CACGACTAGT	5880
TTCCATCATA AATGGTCAA TCAAGTGAGC AACTACACGA ATGCTGGCTG CCAAGTGGCT	5940
CATGACACTT GCCAATTGGT CACGAAGAGC TTCACTCTTG GCCAAGACCC ATGGTGGGGT	6000
CTCATCGATG TATTTATTGG TACGAGAGAT CAGAGTCCAG ACTGCTTCAA GCGCACGTGG	6060
ATAGTCAACT GCTTCCATGT GTGTATGGAA GTCTGCGATT GATTGTWCTG CAACCTCAGC	6120
AAGAACATGA TCATATTCA CGTACACCTTC TACATAGGCA GGGATTTGTC CATCAAAGTA	6180

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CTTATTAATC ATGGAAACCG TACGGTTAAG GAGGTTCCA AGGTCAATTAG CCAATTATA	6240
GTTGATACGG CCGACATAGT CTTCAGGAGT AAAGGTTCCG TCTGAACCAA CTGGAAAGGTT	6300
ACGCATGAGG TAGTAACGAA GTGGATCTAG TCCATAACGC TCTACCAACA TTTCAGGGTA	6360
AACGACATTC CCTTTGACT TAGACATTT TCCGCTTTC ATGACAAACC AACCATGGGC	6420
AATCAAACGA TCAGGTAATT TAACATCCAA CATCATAAGA AGGATTGCC AGTAGATAGA	6480
GTGGAAGCGA AGGATATCTT TTCCTACCAT ATGGAAGACT GTTCCATTCC AGAACTTGTC	6540
AAAGTTACCA TGTTCGTCTT GAGCGTAGCC AAGAGCTGTC GCATAGTTAA GAAGGGCATC	6600
AATCCAAACG TAGACAACGT GTTTGGATT TGATGGGACA GGCACCTCCCC ATGTAAAGGT	6660
TGTACGAGAT ACCGCCAAT CTTCCAAGCC TGGCTCGATG AAGTTGCCGA GCATTTCATT	6720
AAGGCGACCA TCTGGCGTGA TAAATTCAAGG ATGAGCTTG AAAATTCCGA CCAAACGGTC	6780
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TGATGGAGCA ATACCACCG TCACATTTCC AGCTTCATCA CGGAAAACCT CTGCCAGCTG	6900
GCTTCTGTAA AAGAATTCTT CGTCTGATAC TGAATACCAA CCAGAGTATT CACCCAAGTA	6960
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ATCAGTTGTA CGGATAAATT TATCGTATGA GATATCTAGT AATTGCCAGA GTTCTTTAAC	7080
TCCAACCGCC ATTCCATCAA CATAGGCTTG AGGTGTAATA CCAGCTTCTT CCGCTTCTG	7140
CTGGATTTTC TGACCATGTT CATCAAGACC TGTCAGATAA AATACATCGT ACCCCNTCA	7200
GCCTTTGTAA CGTGCTAGGA CATCACATGC GATAGTTGTG TAGGCAGAAC CGATATGAAG	7260
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TCCCTTCCAG GCAAATGAAA CCTGTTTTTC TAACACTTC TTATATCACA TTTTTAATGA	7380
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TCCAACCTCT ATACCTTCTA TTAAACCTTG CAGTAACAAA CCTGCTAAGg CAGTTAAACC	7620
ACTTGCAATT GTCAAGCCAA TTAAGCCACC TAACAAGGTC AAAATCATGG ATTCAATCAA	7680
AAACTGAATT AAAATATTGG CACGTGTTGC ACCCAAAGCC TTACGAAGAC CAATCTCACG	7740
AGTGCCTCT GTCACCGAAA CCAGCATGAT GTTCATGACA CCAGTTCTC CAACAAAGAG	7800
AGAAATCCCT GCGATGGAAC TAATAATCGT CGTCATAAAA CTAAACGATT GTTGAATTTC	7860
TGCAAATACA ACGGACTCAT CTGCCACCTG GTATTCTCCC TGTTGTAAGC CTGCAAGCTC	7920

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TGTCACTTTT CGTGCCAGTT CTGGACCCAG AGTTGGGTT AACTGGTAT CATTCACTCG	7980
AAAGACAATA TTAGCTATT CATCTACATT AAAATTGC A GCAAGGGAGA TATTGGTAGT	8040
AATAGGCAAG CCACCAAACC CATATATT TT TGATCTTTA GCCTCCGGAC TAGTATAAAC	8100
CCCAATGACC CGGTAACCAA ATCCATTGAC TTCTACAACC TTGTTAACAG CCTCTTGAGG	8160
AGATTCAAAT AACTAATGG ACAATTCCCTC ATCTAGCAAA ATGACACTTG CAAACTCTT	8220
GAAATCTTGC TCTCTCAGAC TACGACCTGC AATAATTCA TTCTAACAG CGTCCATGTA	8280
AGTTCTGTTT CCACCTGTCA ATTAGCATT CTCAACCTTT TTATCTTGAT AGGTCAAGAT	8340
GGCATTGTTT GAATTGGTTA CATACTAACT ATCCACTCCC TTCAGTTAG CTGCCTTTG	8400
GACCCAGGAT TCTTGGGTT AACAGGAAC TCCCTTCCTC TTCCAGAAC	8460
CGTAAAAGCT GATTGTTCT GAGTAAAAGA CCCGTCCTTA CTTTTTTAG GAGAGAAAAA	8520
GACGCTAATA TTTTCTGAG ATTTAGTCAT ATCTTTATTG ACTTGACGAG ATAGGGAAATC	8580
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CAAAGAACGC ATCTTGAG CCATGATAGA TGAAAAGCA AATTCAGAT TCTGCATCTT	8700
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TCTGACGTTT GGCATAGGCA GCAATCTCAG GCTCATGCGT TACCATGATA ATGGTTTTTC	8820
CTTCTTTATT CAAATCAACC AATAATTGCA TAATTTGGTT ACCTGTTTG GTATCCAAGG	8880
CTCCGTGCGG TTCATCCGCT AGGATAATAG AAGGATTGTT TACCAAGGCA CGCCCAATGG	8940
CTACACGGTG CTTTGACCA CCAGATAATT CTGAAGGTAA ATGGTGACTA CGTTCTGTCA	9000
ATTCAACCTT GTCTAAATAT TCCTCAGCCA ACTTGCGACG TTTTGAAGAC GAAACTCCTG	9060
CGTAAATCAA GGGCAATTCT ACATTTGCA GAGCATTGAG CTTCGATAGA AGAAAAGACT	9120
GCTGAAAGAC AAAACCGATT TGTTGGTTAC GGACCTTAGC TAGTTGTTT TCACCAAGCC	9180
CAGCCACTTC TTGACCTTCA AGATAATATT CTCCACTGGT TGGTGTATCC AACATGCCAA	9240
TCGTATTCA CAGAGTGGAC TTACCAAGACC CAGATGGTCC CATGATGGCT ACAAAATTCAC	9300
CCTCATTCACTC TTCTAGATTG ATATTTTGAA GAACCTGCAAG TTCTGGTCA CCATTACGGT	9360
AACTCTGAA GATATTTTT AGACTAATTA GTTGCTTCAT CAGCCTTCAC CTCTTTCCCT	9420
TCTTCCAAGG AAGATGTTGG ATTACTGATG ACCTTAGCAC CGTCGTTAA ACCAGAAAGTG	9480
ATTTCTTGAT TTTCTGCGTC AGCATTCCC AATGAAACCT CAACTTTTT AGCCTTTGT	9540
TGTTCATCCA CAATCCAGAC ATAATTTTA CTATCATCCA TTACTAGACT GCTAACAGGA	9600
ACAAGAAATAG CCTTAGTTTT GCTTTAACCC TCAATGTTGA CAGAAAACC TTGTTCAA	9660
TCACCAACCT CGCCTGTCAC ATCAATAGTA TAAGGGTATT TAGAACCTGT ATTATTCCCG	9720

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GCTGCTGGAC TAGCTGCTTC ACCATTGTTT TTAGGATAGT CAGAAATATA GCTTAATTTC	9780
CCAGTCCATT TTTTATCAGG ATACACTTTA GAAGTAAAGC TTACTTCTTG ACCTACAGAA	9840
AGGTTGGCTA GATTGTACTC AGACAATTCT CCCTTGACTT GTAAATTTTC ATTGCTGACA	9900
ATATGAACCA TAACTTGACT CGCCCCTGTT GGAGATTTAG AAACATTGCT ATTGACTTCG	9960
ACCACAGTTC CCTCTAGGGT ACTGAGAACCA GTTGTGCA CCAATTGACT TTGAGCCTTG	10020
CTTAATTGCG CCGCAGCATC TGCACGCGCA TCACGGGCAT CACCCAATTG AGCGTCAATA	10080
GAAGCAACAG AATTTCCAGC CACTGGAGTT GGGCTTTGCA CCGTTGCATC TTCTCCTCCT	10140
ACTGGCGCTG GTAACGTGAG AGCCGGAGCT GAAGCGGCTT CATTGCTGC TTGATTGAGT	10200
TCATTGATAT GACGATCTGC CCTAGCTACT GCTCGACTAG CTGAATCATA GGCGCCTGC	10260
GCTTCTGAAC TACTGTACTT GACTAAAGCC TGCCCTTCGC TGACCTTATC GCCCACAGAA	10320
ACAAGGATTT CATCTAAATC ACCCTTACTA GCATCAAAAT AAACATATTG TTCATTTTT	10380
GCTGTTACTG TCCCTGACAA TAAAACAGAG GAGGCCACGC TTCCCTTCCTT GGCAACAAACA	10440
AGATGAGTAG GCTCATCTTT TAGAGCAGTC TGAGAAGGTT GTCTAAAGAG TAAAATCCCC	10500
CCAGCACCCA ATACAACATAC ACTCGCAGCA CCGATTGCTG CATAACAGTTG CCACCTTTA	10560
GCTTTACCAT TCTTTTCTT CATAATGAAA CTCCCTTCCTT TTTTTACAAT ACTTTGCTAT	10620
TATACCAAAT TTCCCTCCAG CAAACAATAC AGTTCAAGGAT TAAACAATCG TTCGGAATT	10680
TGCTTTTCGG	10690

(2) INFORMATION FOR SEQ ID NO: 94:

- (i) SEQUENCE CHARACTERISTICS:
- (A) LENGTH: 8195 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 94:

GAGAAAGCGC CCACGTTCC CCCAAGGGAG AAAGGCGGAC AGGTATCCGG TAAGCGGCCA	60
GGGTCGGAAC AGGAGAGCGC AACGAGGGAG CTTCCCGAGG GGAAACGCC GGTATCTTTA	120
TAGTCCTGTC GGGTTTCGCC ACCTCTGACT TGAGCGTCGA TTTTTGTGAT GCTCGTCAGG	180
GGGGCGGAGC CTATGGAAAA ACGCCAGCAA CGCGGCCCTT TTACGGTTCC TGGCCTTTG	240
CTGGCCTTT GCTCACATGT TCTTCCCTGC GTTATCCCCT GATTCTGTGG ATAACCGTAT	300
TACCGCCTTT GAGTGAGCTG ATACCGCTCG CCGCAGCCGA ACGACCGAGC GCAGCGAGTC	360

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AGTGAGCGAG	GAAGCGGAAG	AGCGCCCAAT	ACGCAAACCG	CCTCTCCCCG	CGCGTTGGCC	420
GATTCACTAA	TGCAGCTGGC	ACGACAGGTT	TCCCGACTGG	AAAGCGGGCA	GTGAGCGCAA	480
CGCAATTAAT	GTGAGTTAGC	TCACTCATTA	GGCACCCAG	GCTTTACACT	TTATGCTTCC	540
GGCTCGTATG	TTGTGTGGAA	TTGTGAGCGG	ATAACAATT	CACACAGGAA	ACAGCTATGA	600
CaTGATTACG	AATTGAGCT	CGGTACCCGG	AAAATCCAGA	AAATGCTTGA	AAAAATCCT	660
AGAAGATGGT	ATAATACTAA	ATTGTAAGGG	TTATCACATA	TAACCTAAAA	AAAGAAAGAA	720
CAAAAGGAGA	GTCAAACTAT	GGCTTCTAAA	GATTTCCACG	TAGTGGCAGA	AACAGGTATT	780
CACGCACGTC	CAGCAACATT	GTTGGTACAA	ACTGCTAGCA	AATTTGCTTC	AGATATCACT	840
CTTGAGTACA	AAGGTAATC	AGTTAACCTT	AAATCAATT	TGGGTGTTAT	GAGTCTTGGT	900
GTTGCCAAG	GTGCTGACGT	AACTATCTCA	GCTGAAGGTG	CAGATGCAGA	TGACGCTATC	960
GCTGCAATCT	CAGAAACAAT	GGAAAAAGAA	GGATTGCCAT	AAGGGAAATG	ACAGAAATGC	1020
TTAAAGGAAT	CGCAGCATCT	GACGGTGTG	CAGTTGCAA	AGCATATCTA	CTCGTTCAGC	1080
CGGATTGTC	ATTGAGACT	ATTACAGTCG	AAGATACAAA	CGCAGAAGAA	GCTCGCCTTG	1140
ATGCCGCTCT	ACAGGCATCA	CAAGACGAGC	TTTCTGTTAT	TCGCGAGAAA	CGAGTAGGTA	1200
CGCTCGGTGA	AGAACGAGCT	CAAGTTTTG	ATGCTCACTT	AATGGTTCTT	GCTGACCCAG	1260
AAATGATCAG	CCAAATCAAG	GAAACTATCC	GTGCGAAGAA	AGTGAATGCA	GAAGCAGGTC	1320
TGAAAGAAGT	TACAGATATG	TTTATCACTA	TCTTTGAAGG	CATGGAAGAC	AACCCATACA	1380
TGCAAGAACG	CGCAGCGGAT	WTCCCGGACG	TGACAAAACG	TGTATTGGCA	AACCTTCTTG	1440
GTAAAAAATT	GCCAAACCCA	GCTTCTATCA	ATGAAGAAGT	GATTGTGATT	GCGCATGACT	1500
TGACTCCCTC	AGATACAGCT	CAATTGGACA	AAAACCTTGT	AAAAGCTTTT	GTAACCAACA	1560
TTGGTGGACG	TACAAGCCAC	TCAGCTATCA	TGGCACGTAC	ACTTGAAATT	GCTGCTGTAT	1620
TAGGTACAAA	TAACATCACT	GAAATCGTTA	AAGACGGTGA	CATCCTTGCT	TTAACCGGGA	1680
TCACTGGAGA	AGTGATTATC	AACCAACAG	ATGAACAAGC	GGCAGAATT	AAAGCAGCTG	1740
GTGAAGCCTA	TGCGAAACAA	AAAGCTGAAT	GGGCACTTTT	AAAAGATGCT	CAAACAGTGA	1800
CTGCTGACGG	AAACACCTTC	GAGTTGGCTG	CTAATATCGG	TACTCCAAA	GACCTTGAG	1860
GTGTTAACAA	CAACGGTGCA	GAAGCTGTTG	GACTTTACCG	TACAGAGTTTC	TTGTACATGG	1920
ATTCTCAAGA	CTTCCCAACT	GAAGATGAGC	AGTATGAAGC	ATACAAGGCT	GTTCTTGAG	1980
GAATGAACGG	AAAACCTGTT	GTCGTTCGTA	CAATGGATAT	CGGTGGAGAT	AAGGAACCTTC	2040
CTTACTTCGA	TATGCCCTCAC	GAAATGAACC	CATTCCPTGG	ATTCCGTGCT	CTTCGTATCT	2100
CTATCTCTGA	GACTGGAGAT	GCTATGTTCC	GCACACAAAT	CCGTGCTCTT	CTTCGTGCGT	2160

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CTGTTCACGG TCAATTGCGT ATCATGTTCC CAATGGTTGC GCTCTTGAAA GAATTCCGTG	2220
CAGCGAAAGC AGTCCTTGAT GAAGAAAAAG CAAACCTTCT TGCTGAAGGT GTTGCAGTTG	2280
CGGATAACAT CCAAGTTGGT ATCATGATCG AGATTCTGC AGCGGCTATG CTTGCAGACC	2340
AATTGCTAA AGAAGTTGAC TTCTTCTCAA TTGGTACAAA CGACTTGATC CAATATACAA	2400
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TCCTACGCTT GATTAACAAT GTGATCAAAG CAGCTCACCG TGAAAGTAAA TGGGCTGGTA	2520
TGTGTGGTGA GATGGCTGGT GACCAACAAG CTGTTCCACT TCTTGTGGA ATGGGCTTGG	2580
ATGAGTTCTC TATGTCAGCA ACATCTGTAC TTCGTACACG CAGCTTGATG AAGAAACTCG	2640
ACACAGCTAA GATGGAAGAG TACGCAAACC GTGCCCTTAC AGAATGCTCA ACAATGGAAG	2700
AAGTTCTTGA ACTTCAAAAA GAATACGTTA ATTTTGATTA ATCGAAAAGT CCCTGCAACT	2760
CAGTTACAGG GATTTTTTTG ATATTTTAAA AAGAATTTTC AAGAAAATCT TTCTTATAGA	2820
AAGTCCAACC TTGAAAAAGT AGTGGTCAGA ACAAAAATA CTTAAATGGT TCATAAAATT	2880
CTTGACAAGT TGGATATTTA GGAGTAAACT ATTAACCAAGT TAAGTAATAG AGAGGAGTTT	2940
CTGCAATTAA GAAATGAATT GCAACTAGAA ATATCAAATA GAAAGAGAGT TTGATGAAA	3000
ATTAATAAAGA AATACCTTGT TGGTTCTGCG GCACCTTGAT TTTAAGTGT TGTTCTTACG	3060
AGTPTGGACT GTATCAAGCT AGAACGGTTA AGGAAAATAA TCGTGTTC TATATAGATG	3120
GAAAACAAGC GACGCAAAAA ACGGAGAATT TGACTCCTGA TGAGGTTAGC AAGCGTGAAC	3180
GAATCAATGC TGAGCAAATC GTCATCAAGA TAACAGACCA AGGCTATGTC ACTTCACATG	3240
GCGACCACTA TCATTATTAC AATGGTAAGG TTCCCTTATGA CGCTATCATC AGTGAAGAAT	3300
TACTCATGAA AGATCCAAAC TATAAGCTAA AAGATGAGGA TATTGTTAAT GAGGTCAAGG	3360
GTGGATATGT TATCAAGGTA GATGGAAAAT ACTATGTTA CCTTAAGGAT GCTGCCACG	3420
CGGATAACGT CCGTACAAAA GAGGAAATCA ATCGACAAAA ACAAGAGCAT AGTCAACATC	3480
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ATACTACAGA TGATGGTTAT ATCTTTAATG CTTCTGATAT CATAGAGGAT ACTGGTGATG	3600
CTTATATCGT TCCTCATGGA GATCATTACC ATTACATTCC TAAGAATGAG TTATCAGCTA	3660
GCGAGTTGGC TGCTGCAGAA GCCTTCCTAT CTGGTCGAGG AAATCTGTCA AATTCAAGAA	3720
CCTATCGCCG ACAAAATAGC GATAACACTT CAAGAACAAA CTGGGTACCT TCTGTAAGCA	3780
ATCCAGGAAC TACAAATACT AACACAAGCA ACAACAGCAA CACTAACAGT CAAGCAAGTC	3840
AAAGTAATGA CATTGATAGT CTCTTGAAAC AGCTCTACAA ACTGCCTTG AGTCAACGAC	3900

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GTGTTGCAGT	GCCACACGGA	GATCATTACC	ACTTCATCCC	TTACTCTCAA	ATGCTGAAT	4020	
TGGAAGAACG	AATCGCTCGT	ATTATTCCCC	TTCGTTATCG	TTCAAAACCAT	TGGGTACCA	4080	
ATTCAAGGCC	AGAACAAACCA	AGTCCACAAAC	CGACTCCGGA	ACCTAGTCCA	GGCCCGCAAC	4140	
CTGCACCAAA	TCTTAAAATA	GACTCAAATT	CTTCTTTGGT	TAGTCAGCTG	GTACGAAAAG	4200	
TTGGGGAAGG	ATATGTATT	TC GAAGAAAAGG	GCATCTCTCG	TTATGTCTTT	GCGAAAGATT	4260	
TACCATCTGA	AACTGTTAAA	AATCTGAAA	GCAAGTTATC	AAAACAAGAG	AGTGTTCAC	4320	
ACAC	TTAAC	TGCTAAAAAA	GAAAATGTTG	CTCCTCGTGA	CCAAGAATT	4380	
CATATAATCT	GTAA	CTGAG	GCTCATAAAG	CCTTGTGAA	AAATAAGGGT	4440	
ATTTCCAAGC	CTTAGACAAA	TTATTAGAAC	GCTTGAATGA	TGAATCGACT	AAATAAGAAA	4500	
AATTGGTAGA	TGATTTAT	GCATTCTAG	CACCAATTAC	CCATCCAGAG	CGACTTGGCA	4560	
AACCAATT	TC	AAATTGAG	TATACTGAAG	ACGAAGTTCG	TATTGCTCAA	4620	
AGTATACAAC	GTCAGATGGT	TACATTTTG	ATGAACATGA	TATAATCAGT	GATGAAGGAG	4680	
ATGCATATGT	AA	GCCTCAT	ATGGCCATA	GTCACTGGAT	TGGAAAAGAT	4740	
ATAAGGAAA	AGT	TGCA	CTAAAGCTATA	CTAAAGAAA	AGGTATCCTA	4800	
CAGACGCAGA	TGTTAAAGCA	AATCCAAC	TG GAGATGTC	AGCAGCTATT	TACAATCGTG	4860	
TGAAAGGG	AA	ACGAATT	CCACTCGTC	GACTTCCATA	TATGGTGAG	4920	
AGGTTAAA	AA	CGGTAATT	TG ATTTCCTC	ATAAGGATCA	TTACCATAAT	4980	
CTTGGTTG	TGAT	CACACA	TACAAAGCTC	CAAATGGCTA	TACCTTGAA	5040	
CGACGATTAA	GTACTACG	TA	GAACACCC	TC	ACATTCTAA	5100	
GCAATGCCAG	TGAGCATG	TG	TTAGGCAAGA	AA	GACCAAG	5160	
TCAAAGCGGA	TGAAGAGCCA	GTAGAGGAA	CA	CTGCTGA	CCCAGAA	5220	
AGACTGAAA	AGT	AGAAGCC	CAACTCAA	AG	AGCAGAA	5280	
ATTCTAGTCT	GAAAGCCA	AT	GCAACAGAA	CTCTAGCTG	TTTACGAA	5340	
TTCAAATTAT	GGATAACAA	AT	AGTATCATG	CAGA	AGCAGA	5400	
AAGGAAGTAA	TCTT	CATCT	GTAAGTAAG	AAA	AACTAA	5460	
CGATAAAAGAG	GCTT	CATT	TTATTATG	TATATG	AA	TTCTGACA	5520
AAAAGAGTAA	ACTATTAA	ACT	AGTAA	CCGG	TTATTAGT	5580	
TACTTAAGAA	AAGAGGAA	AG	AATGAA	AA	ATCTAGCAG	5640	
GTCCTTGCCC	TAAGTGT	TTG	TTCCTATGAA	CTTGGT	CGTC	ACCAAGCTGG	
						TCAGGTTAAG	
						5700	

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AAAGAGTCTA ATCGAGTTKC TTATATAGAT GGTGATCAGG CTGGTCAAA GGCAGAAAAC	5760
TTGACACCAG ATGAAGTCAG TAAGAGGGAG GGGATCAACG CGAACAAAT CGTCATCAAG	5820
ATTACGGATC AAGGTTATGT GACCTCTCAT GGAGACCATT ATCATTACTA TAATGCCAG	5880
GTCCTTATG ATGCCATCAT CAGTGAAGAG CTCCATGAA AGATCCGAA TTATCAGTG	5940
AAGGATTCAAG ACATTGTCAA TGAAATCAAG GGTGGTTATG TTATCAAGGT AGATGGAAA	6000
TACTATGTTT ACCTTAAGGA TGCAGCTCAT GCGGATAATA TTCGGACAAA AGAAGAGATT	6060
AAACGTCAGA AGCAGGAACA CAGTCATAAT CACGGGGGTG GTTCTAACGA TCAAGCAGTA	6120
GTTGCAGCCA GAGCCAAGG ACGCTATACA ACGGATGATG GTTATATCCT CAATGCATCT	6180
GATATCATTG AGGACACGGG TGATGCTTAT ATCGTTCTC ACGGCGACCA TTACCAATTAC	6240
ATTCCTAAGA ATGAGTTATC AGCTAGCGAG TTAGCTGCTG CAGAAGCCTA TTGGAATGGG	6300
AAGCAGGGAT CTCCGCTCTC TTCAAGTTCT AGTTATAATG CAAATCCAGC TCAACCAAGA	6360
TTGTCAGAGA ACCACAATCT GACTGTCACT CCAACTTATC ATCAAATCA AGGGAAAAC	6420
ATTTCAAGCC TTTTACGTGA ATTGTATGCT AAACCCATT CAGAACGCCA TGTGGAATCT	6480
GATGCCCTTA TTTTCGACCC AGCGCAAATC ACAAGTCGAA CCGCCAGAGG TGTAGCTGTC	6540
CCTCATGGTA ACCATTACCA CTTTATCCCT TATGAACAAA TGTCTGAATT GGAAAACGA	6600
ATTGCTCGTA TTATTCCCCT TCGTTATCGT TCAAACCATT GGGTACCCAGA TTCAAGACCA	6660
GAACACCAA GTCCACAATC GACTCCGGAA CCTAGTCCAA GTCCGCAACC TGCACCAAT	6720
CCTCAACCAAG CTCAGCAA TCCAATTGAT GAGAAATTGG TCAAAGAACG TGTCGAAAA	6780
GTAGGCGATG GTTATGCTT TGAGGAGAAT GGAGTTCTC GTTATATCCC AGCCAAGGAT	6840
CTTTCAAGCAG AACACAGCAGC AGGCATTGAT AGCAAACGG CCAAGCAGGA AAGTTTATCT	6900
CATAAGCTAG GAGCTAAGAA AACTGACCTC CCATCTAGTG ATCGAGAATT TTACAATAAG	6960
GCTTATGACT TACTAGCAAG AATTCAACCA GATTACTTG ATAATAAAGG TCGACAAGTT	7020
GATTTTGAGG CTTTGGATAA CCTGTTGGAA CGACTCAAGG ATGTCyCAAG TGATAAAAGTC	7080
AAGTTAGTGG ATGATATTCT TGCCTCTTA GCTCCGATTC GTCATCCAGA ACGTTTAGGA	7140
AAACCAAATG CGCAAATTAC CTACACTGAT GATGAGATTG AAGTAGCCAA GTTGGCAGGC	7200
AAGTACACAA CAGAAGACGG TTATATCTT GATCCTCGTG ATATAACCAG TGATGAGGGG	7260
GATGCCCTATG TAACTCCACA TATGACCCAT AGCCACTGGA TTAAGGAGA TAGTTTGCT	7320
GAAGCTGAGA GAGCGGCAGC CCAGGCTTAT GCTAAAGAGA AAGGTTTGAC CCCTCCCTCG	7380
ACAGACCATC AGGATTCAAGG AAATACTGAG GCAAAAGGAG CAGAACGCTAT CTACAACCGC	7440

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GTGAAAGCAG CTAAGAACGT GCCACTTGAT CGTATGCCTT ACAATCTTCA ATATACTGTA	7500
GAAGTCAAA ACGGTAGTTT AATCATACCT CATTATGACC ATTACCCATAA CATCAAATTT	7560
GAGTGGTTG ACGAAGGCCT TTATGAGGCA CCTAAGGGGT ATAATCTTGA GGATCTTTG	7620
GCGACTGTCA AGTACTATGT CGAACATCCA AACGAACGTC CGCATTAGA TAATGGTTT	7680
GGTAACGCTA GCGACCAGT TCGTAAAAAT AAGGTAGACC AAGACAGTAA ACCTGATGAA	7740
GATAAGGAAC ATGATGAAGT AAGTGAGCA ACTCACCTG AATCTGATGA AAAAGAGAAT	7800
CACCGCTGGTT TAAATCCTTC AGCAGATAAT CTTTATAAAC CAAGCACTGA TACGGAAGAG	7860
ACAGAGGAAG AAGCTGAAGA TACCACAGAT GAGGCTGAAA TTCCCTCAAGT AGAGAATTCT	7920
GTATTAAACG CTAAGATAGC AGATGCGGAG GCCTTGCTAG AAAAGTAAC AGATCCTAGT	7980
ATTAGACAAA ATGCTATGGA GACATTGACT GGTCTAAAAA GTAGTCTTCT TCTCGGAACG	8040
AAAGATAATA ACACATTTTC AGCAGAAGTA GATAGTCTCT TGGCTTTGTT AAAAGAAAGT	8100
CAACCGGCTC CTATACAGTA GTAAAATGAA TGGAGCATAT TTTATGGAGA AGTAACCTT	8160
CGTGTACTT CTCTTTTTA GAAAAACGTA ACAGA	8195

(2) INFORMATION FOR SEQ ID NO: 95:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 2004 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 95:

TTTACTAAA GGAAAAAAGA ACTGATTCT CAGTCCTTCA TTAATCTTAT TCCACACTAA	60
ATAGGTATGG GTAAACAGGT TGTTGACCTT GGTGAATCTC GACTTCAACG TCTTCGAATT	120
CTTCTACGAT TTCTTGAGCG ATTTCATTTG CAAAGTCTTC GCTTCCGTCT TCACCTACAT	180
AGAAGGTTAC GATTTCACTG TCTTCATCCA ACATATGTTT CAAGGTTCA GTCAATGTTT	240
GGTGCAATAC AGGGTTTGAC ACAAGAATT TTCCATCCAC CATACTAAA TTATGTTTT	300
CATGGATTTC TAAGCCATCG ATCGTTGAT CACGCACGGC TGTTGTGACG CTTCCGCTAA	360
CGACATCGCT AAGAGCAGCT GTCATACCGCT CTTGGTTTTC TTCAATGGAC TTGCTTGGAT	420
CAAAGGCAAG AAGACTTGTC ATACCTTGAG GAAGAGTGCG AGCCTCTACC ACTACCGCTG	480
GTTGCTCCAA AACTCTGCC GCAGATTGAG CTGCCATGAA GATGTTCTG TTGTTGGCA	540
AGAAGATGAT GTTACGGCA TTAACCTGTT CAACAGCCTT GATAAAGTCT TCTGTTGAAG	600
GGTTCATGGT TTGACCGCCT TCGATAAACAT AATCCACGCC TTGAGAACAG AAGATATCTG	660

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CTAGACCTTT ACCAGCCACC ACAGCAATCA AAGCATACTC TTTTCTTCA GCCGACTTGA	720
TAACCTGAGT AGCTCTTTC TCAACCTGTG CTTCGTGTTG GTTACGCATA TTGTCAACTT	780
TTACCTTGAC CAAGCTACCA TATTTGAGAC CTTCTTGAT AACAAAGCTT GGATCTTCTG	840
TATGAACATG GACTTTGACA ATTTCATCAT CGTTAACAAAC AAGGAGAGAA TCTCCAAGCT	900
CATCCAAGTA GTTACGGAAT TCATCGTAGT CAAAATCTT AGCATAGGTT GGACCTTGCT	960
TAAGAGCTAC CATGATTCA GTACAGTAAC CAAACGTGAT GTCTCTGAC GCTACGTGAC	1020
CAGCTACAGA CTTATGATGC TCTACATTGA TCATCTCACT CATGTTGGCA GGAGTCGCTA	1080
CAAAGTCCTC AGATGCAATA TATTCGCCAG TAAGGGCTGA AAGGAAACCT TCGTAGATGA	1140
AGACCAATCC TTGACCACCT GAGTCCACAA CGCCAACCTC TTTCAATACT GGAAGCATGT	1200
CTGGTGTMTT AGCTAGAGCT GTTTTAGCAC CTTCCAAGGC TGCGCGCATG ACTTCAACAG	1260
CGTCATCTGT TTGCTCAGCT TTTTCTTAG CACCGATAGC AGCTCCACGA GAAACTGTTA	1320
AAATCGTTCC TTCACACAGGT TTCATCACTG CCTTATAGGC AACTTCCACA CCTGATTGGA	1380
AGGCCAGAGC CAAGTCTTGA CCTGTTAACT CGTCTTATAC CTTGATAGCT TGGGAAAATC	1440
CACGGAAAAG CTGAGACGTA ATCACTCCTG AGTTCCCACG CGCACCCATC AAAAGCCCTT	1500
TGGCAAGAAT GCTCGCTACT TCTCCAAGT TAGAAGCTGG CTTGCTGCA ACTTCTTTAG	1560
CACCATTTTC AATGGTCATT CCCATATTTG TCCCAGTATC TCCATCTGGA ACTGGAAAGA	1620
CGTTTAATGA ATTGACATAT TCAGCTTGCT TATTCAAGGC AGTTGATGCA GCCTGCCACCA	1680
TTTCTTGAAA TAAGCTAGTA GTAATTGGT ACACGGTTAT TCTCCTACAA CTTTGATATT	1740
TTGAATGTAG ACATTTACAG TCTGAGGAGT AATTCCAAGC TGTTTTCCA AGCTAAAGGC	1800
AACACGCTCT TGAATGTTT TTGACACTTC ACTAATCTT GTTCCGTAGC TTAACACGGT	1860
ATATACATCA ACTGCAATAC TGCCATCTTC GGCTGCCTT ACGACGACAC CTTTAGAATA	1920
ATTTCCTTA CCTAGCAGGG CTTGGAAATT ATCTTGAGG GCATTTTAC TAGCCATACC	1980
GACCACACCA GAAATCTCAG TTGC	2004

(2) INFORMATION FOR SEQ ID NO: 96:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 11915 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 96:

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CGGGGTTGGG	CTGTTGCC	ATTAAAGCGG	CACCAAGCT	GGGTCAGAA	CGTCGTGAGA	60	
CAGTTCGGTC	CCTATCGTC	GCGGGCGTAG	GAAATTGAG	AGGATCTGCT	CCTAGTACGA	120	
GAGGACCAGA	GTGGACTTAC	CGCTGGTGT	CCAGTTGTCT	TGCCAAAGGC	ATCGCTGGGT	180	
AGCTATGTA	GGAAGGGATA	AACGCTGAAA	GCATCTAAGT	GTGAAACCCA	CCTCAAGATG	240	
AGATTTCCTA	TGATTATATA	TCAGTAAGAG	CCCTGAGAGA	TGATCAGGTA	GATAGGTTAG	300	
AAGTGGAA	GTGGCGACAC	ATGTAGCGGA	CTAATACTAA	TAGCTCGAGG	ACTTATCCAA	360	
AGTAAC	TGAG	AATATGAAAG	CGAACGGTT	TCTTAAATTG	AATAGATATT	CAATTGAG	420
TAGGTATTAC	TCAGAGTTAA	GTGACGATAG	CCTAGGAGAT	ACACCTGTAC	CCATGCCGAA	480	
CACAGAA	CTA	AAGCCCTAGA	ACGCCGGAAG	TAGITGGGGG	TTGCCCTG	TGAGATAGGG	540
AAGTCGCTTA	GCTCTAGGGA	GTTTAGCTCA	GCTGGGAGAG	CATCTGCC	ACAAGCAGAG	600	
GGTCAGCGGT	TCGATCCC	GT	TAAC	TTAGGGGT	GTAGTTAGT	GGTAAACTA	660
CAGCCTTCCA	AGCTGTTGTC	GGCAGTTCGA	TTCTCGTCAC	CCGCTTGAA	CTTTGTTCTT	720	
TGTACCAAGT	TTTGACTTG	GGCGCGTAGC	TCAGGTGGTT	AGAGCGCACG	CCTGATAAGC	780	
GTGAGGTCGG	TGGTTCGAGT	CCACTCGTGC	CCATAGTGT	TAGTCATTA	CTAGGGGATT	840	
GGAATATTAT	CTGTTCACTA	AGAGGACACG	GGCTTGTTCC	CGTATAAACT	ATTTGGAGG	900	
ATTACCAAG	TCCGGCTGAA	GGGAACGGTC	TTGAAAACCG	TCAGCGTGT	AAAAGCGTGC	960	
GTGGGTTCGA	ATCCCACATC	CTCCTTTAT	ATTAACCGGG	GATGGAGCAG	CTCGTAGCT	1020	
CGTCGGGCTC	ATAACCCGAA	GGTCGTAGGT	TCAAATCCTG	CTCCCGCAAT	AAGGCTCGGT	1080	
AGCTCAGTTG	GTAGAGCAAT	GGATTGAAGC	TCCATGTGTC	GGCGGTTCGA	TTCCGTC	1140	
CGCCATTAT	ATATTTGGA	AGGGTAGCGA	AGAGGCTAAA	CGCGCGGAC	TGTAAATCCG	1200	
CTCCTTCGGG	TTCGGGGTT	CGAATCCCTC	CCCTTCCATT	TTACGGGCAT	AGTTAAAGG	1260	
TAGAACTAAG	GTCTCCAAA	CCTTCAGTGT	GGGTCATT	CCTACTGCC	GTGTTAATAG	1320	
AATTATGGCG	GGTGTGGTGA	AGTGGTTAAC	ACACCAAGATT	GTGGCTCTGG	CATGCGTGGG	1380	
TTCGATCCCC	ATCACTCGCC	TATTTATAT	TGGGTATAG	CCAAGCGGTA	AGGCAAGGGA	1440	
CTTGACTCC	CTCATGCGTT	GGTCGAATC	CAGCTACCCC	AGTTACTATT	TGCCGGCTG	1500	
GCGGAATTGG	CAGACCGCCT	GGACTCAAA	TCCAGTGTCC	GCAAGGACGT	GGCGGTTCGA	1560	
CCCCGGCCGC	CGGTATAGTA	TAGTGTAGG	AACGTTGTTA	TTCTCGTTC	CTTTTTATA	1620	
TTATTTTGG	TATAATTATA	GTTATTCAAA	TTTTATTAG	ATTAAGAAAG	TGTAGGGGAG	1680	
TATGTCTTGT	TCTATCGATT	TATTAACACA	TCGGTATTG	AAAAATATTA	AAGAAAATCC	1740	
TGAATTGTTT	GTGCGAATTG	AGTTGGAGTA	TCCTGTTGCA	AGTTTAAAGAG	GGGATGCTAC	1800	

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AGATGTTGAA	GTTATGAAGG	ATCTATTTCA	TTATTTAGTT	TCTACTTTGG	ATCTCACCGT	1860
AGCAAAGGTA	GATGATTTTG	GCAATCTGAT	CCAGTTAGTA	GATCCGATAA	GTCAGGATGC	1920
TATTTTATTT	GAAGTTTCCT	ATACAACGAT	TGAGTTTGCA	TTTGGTAAGG	CTGAAACGAT	1980
TCAAGAGGTC	GAAAATCGTT	TCAATAATT	TATGAATGTA	ATTCAAGAGAA	AGTTAGCTGA	2040
ATCAAATCAT	GCTATTGTTG	GCTGTGGTAT	CCATCCCAC	TGGGATAAAA	ATGAGAATTG	2100
TCCAGTGGCT	TATCCACGCT	ATCAGATGTT	GATGGATTAT	TTGAATTGTA	GTAGAAATAT	2160
TATTAATCA	GATTTACATC	ATTTCCCTGA	ATATGGTACT	TTTATCTGTG	GGAGCCAGGT	2220
TCAGCTGGAT	ATTCAAAAAA	CCAACTACTT	ACGGGTGATT	AATGTTTTA	CTCAAATTGA	2280
AGCGGCTAACG	GCTTATTTAT	TTGCAAACTC	TGAATTTCG	GGTGCAGGATT	GGGATACGAA	2340
AATTCAAGG	GATATTTCT	GGGAAGAAC	TATGCATGGT	ATCTATCCAG	AGAATGTTGG	2400
GGTCAATGCT	AGACTCCTTA	ATGATGAAAC	TGATTTTTT	GACTATCTAA	ATCATTCTGC	2460
GATTTTACT	CGGGAACGTG	ATGGGCAGAC	CTATTATTT	TATCCTATT	AGGCTGGGGA	2520
CTATTGGCT	ACGTCCGAAA	TCCAAGCATT	TGCTCTGAAT	GGGGATGAGG	TTATTATTTA	2580
CCCCAAGAG	AAGGATTTG	AAACTCATCG	TAGTTACCAG	TACCAAGATT	TAACGACTCG	2640
AGGAACAGTT	GAGTTTCGTA	GTGTGTGTAC	ACAGCCACTT	GATAGGACTT	TTGCTTCTGC	2700
AGCTTTCAC	TTGGGATTAT	TGGTTAATT	AGACAAGTTA	GAAGCTTACT	TAGAAACAGC	2760
ACCTTCTTT	AAAGTATTTG	GTTATGATTA	CAAGTCTTTA	AGGAGACAAT	TTTCTAAGAA	2820
AAATCTTACA	GATGAGGAAG	AAACTACGAT	TATTGAATT	TCCAAAGACT	TACTCCTACT	2880
AGCTGAGGAG	GGACTAGTGG	TGAGAAATAA	GGAGGAAATG	ACCTATTTAC	AGCCTTTGAG	2940
AGAAGAATTG	AGCCTATAAT	TTCTCTTATA	AAGGGAGAAT	TTTCTGAAAA	ATCATGATAT	3000
AATGGACGAG	ACTATAGATA	AAGGATAGAG	AGTAATGACA	TTAGTTTATC	AATCAACCGC	3060
TGATGCCAAC	AATACAGTAA	CTGCCAGCCA	AGCAATTTG	CAAGCTTG	CGACGGACGG	3120
CGGTTGTTT	ACACCGGATA	CTTATCCAA	GGTAGATTTG	AACTTGACA	AATTGAAAGA	3180
TGCTTCTTAC	CAGGAAGTTG	CTAAGCTAGT	TTTGTCA	TTTTAGATG	ACTTTACAGT	3240
TGAGGAGTTG	GACTACTGTA	TCAACAATGC	CTACGATAGC	AAATTTGATA	CTCCAGCTAT	3300
TGCACCATTA	GTGAAATTAG	ATGGGCAATA	CAATTGGAA	CTTTCCATG	GTTCAACGAT	3360
TGCCTTTAAG	GATATGGCCT	TGTCTATTT	GCCATACCTT	ATGACGACTG	CTGCTAAGAA	3420
ACATGGTTG	GAGAACAAAGA	TTGTTATCTT	GACAGCGACA	TCTGGTGACA	CGGGAAAGC	3480
TGCTATGGCG	GGGTTTGCAGA	ATGTGCCTGG	TACTGAGATT	ATCGTCTTT	ATCCAAAGGA	3540

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TGGTGTCAAG AAGATTCAAG AGTTACAAAT GACCACTCAAG ACTGGCGACA ATACTCATGT	3600
TATTGCTATT GATGGTAACT TTGACGATGC GCACACAAAT GTGAAGCACA TGTTAACGA	3660
CGTGGCTCTT CGTGAAAAT TGACTACCAA CAAGTTGCAA TTTTCATCAAG CTAACTCTAT	3720
GAACATTGGT CGTCTGGTGC CACAAATTGT TTATTATGTT TATGCTTACG CTCAATTGGT	3780
TAAGACTGGT GAAATTGTAG CTGGTGAAAA GGTTAACTTC ACAGTACCAA CAGGAAACTT	3840
TGGAATATC TTGGCTGCCCT TTATGCCAA ACAATTGGT TTGCCAGMTG GTAAATTAAT	3900
CTGTGCTTCA AATGACAACA ATGTTTGAC AGACTTCTTT AAAACACGTG TCTATGACAA	3960
AAAACGTGAG TTTAAGGTAA CAACCAAGCCC ATCTATGGAT ATCTTGGTAT CTTCAAACCTT	4020
GGAGCCCTTG ATTTTCCATC TTTTGGAAA TAATGCTGAA AAGACAACTG AACTTATGAA	4080
TGCCTTGAAC ACGCAAGGAC AAATAAAGTT GACAGACTTT GATGCAGAGA TTTTGGACCT	4140
CTTTCAGCT GAATATGCGA CTGAGGAAGA AACGGCAGCA GAGATCAAGC GTGTTGTGA	4200
GTTAGATTCT TATATCGAGG ACCCTCATAC AGCTGTTGCT TCAGCAGTTT ATAAAAAATA	4260
CCAATCGGCC ACTGGAGATG TAACTAAGAC AGTGATTGCT TCAACAGCTA GTCCATACAA	4320
GTTCCAGTA GTTGCAGTAG AAGCTGTAAAC TGGAAAAGCA GGTTAAACAG ACTTTGAAGC	4380
CTTGGCTCAA TTACATGAAA TCTCAGGCCT TGCACTGCCA CCAGCAGTTG ATGGGCTTGA	4440
AATAGCTCCA ATTCGTCACA AGACAAACAGT GGCAGCTGCT GACATGCAAG CAGCGGTTGA	4500
GGCTTATTAA GGACTTTAAG ACAGAGGGAG CAAACTCGGT TGGGAAACCA ACTGAGTTTC	4560
TTTTCATCG GAGGAGAGAT TGTTTAAGAA AAATAAAGAC ATTCTTAATA TTGCATTGCC	4620
AGCTATGGGT GAAAATTTC TGCACTGCT AATGGGAATG GTGGACAGTT ATTTGGTTGC	4680
TCATTAGGA TTGATAGCTA TTTCAGGGGT TTCAGTAGCT GGTAATATTA TCACCATTAA	4740
TCAGGGCATT TTCACTCGCTC TGGGAGCTGC TATTTCCAGT GTTATTCAA AAAGCATAGG	4800
GCAGAAAAGAC CAGTCGAAGT TGGCTTATCA TGTGACTGAG GCGTTGAAGA TTACCTTACT	4860
ATTAAGTTTC CTTTTAGGAT TTTTGTCCAT CTTCGCTGGG AAAGAGATGA TAGGACTTTT	4920
GGGGACGGAG AGGGATGTAG CTGAGAGTGG TGGACTGTAT CTATCTTGG TAGGCAGGATC	4980
GATTGTTCTC TTAGGTTAA TGACTAGTCT AGGAGCCTTG ATTCGTGCAA CGCATAATCC	5040
ACGTCTGCCCT CTCTATGTTA GTTCTTATC CAATGCCCTG AATATTCTTT TTTCAAGTCT	5100
AGCTATTTTT GTTCTGGATA TGGGGATAGC TGGTGTGCT TGGGGACAA TTGTGTCTCG	5160
TTTGGTTGGT CTTGTGATTG TGTGGTCACA ATTAAAAGTG CCTTATGGGA AGCCAACTTT	5220
TGGTTTAGAT AAGGAAGTGT TGACCTTGGC TTTACCAAGCA GCTGGAGAGC GACTPATGAT	5280
GAGGGCTGGA GATGTAGTGA TCATTGCCCTT GGTGTTCTT TTTGGGACGG AGGCAGTTGC	5340

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TGGGAATGCA ATCGGAGAAG TCTTGACCCA GTTTAACTAT ATGCCTGCCT TTGGCGTCGC	5400
TACGGCAACG GTCATGCTGT TGGCCCGAGC AGTTGGAGAG GATGATTGGA AAAGAGTTGC	5460
TAGTTTGAGT AAAACAAACCT TTTGGCTTTC TCTGTTCCCTC ATGTTGCCCT TGCTCTTAG	5520
TATATATGTC TTGGGTGTAC CATTAACTCA TCTCTATAACG ACTGATTCTC TAGCGGTGGA	5580
GGCTAGTGTGTT CTAGTGACAC TGTTTCACT ACTTGGGACC CCTATGACGA CAGGAACAGT	5640
CATCTATAACG GCAGTCTGGC AGGGATTAGG AAATGCACGC CTCCCTTTT ATGCGACAAG	5700
TATAGGAATG TGGTGTATCC GCATTGGGAC AGGATATCTG ATGGGGATTG TGCTTGGTTG	5760
GGGCTTGCCCT GGTATTTGGG CAGGGTCTCT CTTGGATAAT GGTTTCGCT GGTTATTCT	5820
ACGCTATCGT TACCAAGCGCT ATATGAGCTT GAAAGGATAG GAAATGCAA AAACAGCTTT	5880
TATTGGGAT TTAGACGGGA CTTTATTGGA CTCTTACGAA GCGATTTTAT CAGGGATTGA	5940
GGAGACTTTT GCTCAGTTTT CTATTCCCTA TGATAAGGAG AAGGTGAGAG AGTTTATCTT	6000
CAAGTATTG TGCAAGATT TGCTTGTGCG GGTGGCAGAA GATAGAAATC TGGATGTTGA	6060
GGTGCTAAAT CAGGTGCGTG CCCAGAGTCT GGCTGAGAAG AATGCTCAGG TAGTTTGAT	6120
GCCAGGTGCG CGTGAGGTGC TAGCTTGGGC ACAGCAATCA GGAATTTCAGC AGTTTATATA	6180
TACTCATAAG GGGAAACAAACG CTTTTACCAT TCTCAAGGAC TTGGGGGTGG AATCCTATTT	6240
TACAGAGATT TTAACCAGTC AGAGTGGCTT TGTGCGGAAG CCAAGTCCAG AAGCGGCTAC	6300
CTATCTGCTA GATAAGTATC AGTTGAATTG TGATAATACT TATTATATAG GGGATCGGAC	6350
TCTGGATGTG GAATTTGCCG AGAATAGTGG GATTCAAAGC ATCAACTTT TAGAGTCTAC	6420
TTATGAGGG AATCACAGGA TTCAAGCGTT AGCAGATATT TCCCCTATT TTGAGACTAA	6480
GTGATAAAAA GATTGTGTCA GTTTGTGAC AGAGACCTAA CAAACTATT CAAGTAACCT	6540
AGTTGTTAC AAGGAATAGA CAGTTCTGTT AAATAGGCCG GAGAGGGCTT TTTTCTACA	6600
TTTTTGTGT TATGATAGAC AGGTACTCAT TTGAAAGGAA TTTGAAAGAA TGAAGAAAAG	6660
AATGTTATTA GCGTCAACAG TAGCCTTGTC ATTTGCCCA GTATTGGCAA CTCAAGCAGA	6720
AGAAGTTCTT TGGACTGCAC GTAGTGTGAA GCAAATCCAA AACGATTGAA CTAAAACGGA	6780
CAACAAAACA AGTTATAACCG TACAGTATGG TGATACTTTG AGCACCATGG CAGAACCTT	6840
GGGTGTAGAT GTCACAGTGC TTGCGAATCT GAACAAAATC ACTAATATGG ACTTGATT	6900
CCCAGAAACT GTTTGACAA CGACTGTCAA TGAAGCAGAA GAAGTAACAG AAGTTGAAAT	6960
CCAAACACCT CAAGCAGACT CTAGTGAAGA AGTGACAACG GCGACAGCAG ATTTGACCA	7020
TAATCAAGTG ACCGTTGATG ATCAAACCTGT TCAGGTTGCA GACCTTCTC AACCAATTGC	7080

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AGAAGTTACA AAGACAGTGA TTGCTTCGCA AGAAGTGGCA CCATCTACCG GCACCTCTGT	7140
CCCAGAGGAG CAAACGACCG AAACAACCTCG CCCAGTTGAA GAAGCAACTC CTCAGGAAAC	7200
GACTCCAGCT GAGAACAGG AAACACAAGC AAGCCCTCAA GCTGCATCAG CAGTGGAACT	7260
AACTACAACA AGTTCAGAAG CAAAAGAACT AGCATCATCA AATGGAGCTA CAGCAGCAGT	7320
TTCTACTTAT CAACCAGAAG AGACGAAAT AATTCAACA ACTTACGAGG CTCCAGCTGC	7380
GCCCCGATTAT GCTGGACTTG CAGTAGCAAA ATCTGAAAT GCAGGTCTTC AACCACAAAC	7440
AGCTGCCTTT AAAGAAGAAA TTGCTAACTT GTTTGGCATT ACATCCTTA CTGGTTATCG	7500
TCCAGGAGAC AGTGGAGATC ACGGAAAAGG TTTGGCTATC GACTTTATGG TACCAAGACG	7560
TTCAAGAATTA GGGGATAAGA TTGCGGAATA TGCTATTCAA AATATGGCCA GCCGTGGCAT	7620
TAGTTACATC ATCTGAAAC AACGTTTCTA TGCTCCATTG GATAGCAAAT ATGGGCCAGC	7680
TAACACTTGG AACCCAATGC CAGACCGTGG TAGTGTGACA GAAAATCACT ATGATCACGT	7740
TCACGTTTCA ATGAATGGAT AAACCCGACT TGATAACATC ATTTTGACGA ATGAGATCTA	7800
GCTTTCGTGA TGGAAAGCGA TTCTCGTTG TTTTTCTTT GTCATACTCT TCGAAAATCT	7860
CTTCAAAACCA CGTCAGTTTT ATCTGAAACT TCAAAGCTGT GCTTGAGCA ACCTGCGACT	7920
AGCTTCCTAG TTTGCTTTT GATTTTCATT GAGTATCAAT TTGAATGGAA AATGGAAAGT	7980
TATCATCTTG TAATGAGTTA AGCAACATTC TTGCAATCTA TTTTACTTTA TATCACAAATT	8040
AATTAGTCAA ATATTGATAA ATCAATAAAA AGAGAGGGGA AGAAATGCTA GAGATTCAAG	8100
ATTTACTGTA TCAACTCCGC TTGCTGAGC AAGCGAGTAC GCAATTGTTT GAAAAAAGGC	8160
TTGGGATTAG TTTGACACGG TATCAGATT TACTGTTTT GCTGGAGCAT TCTCCTTGT	8220
ACCAAATGGC GTTTCAGGAG CGTTGAAAAT TTGATCAGGC TGCTTGACA CGGCATTTC	8280
AAATTTGGA AACCGAAGGT TTGGTGGAGC GTCATCGTAA TCCTGAAAT CAGCGGGAAAG	8340
TGTTGGTAGA GGCTGCGAAG TATGCCAAGG AGCAGTTAGT GGTGAATCCC CCTCTGCAAC	8400
ATATCAGGGT TAAGGAAGAG ATAGAAAGTA TCTTAACAGA GTTTGAGAGA ACAGAACTCA	8460
GCCGTTTATT AAATAAAATTG GTTTGGGTA TTGAAAATAT AGAAATTAA GGAGAAATAG	8520
ATGTCAAATTAA TTTTAACAC GATCGTTGCT TTGGAGCATT TTACATTAA TTATTTGGAA	8580
AGTATTGCCA CGCAATCAGA TGCGACTAGT CGTGTATTAA ATATGGAAA GGAAGAATTG	8640
GCTCATCCGT CAGTAAGTTC ATTGTTCAA AATCAAGGAA TTTATAAGGC TCTGCTAGGA	8700
GTCTTCTCT TGATGTCAT TTATTTCTCA CAGAATTAG AAATTGTCAC TATTTTTGTC	8760
TTATTTGTGA TTGGTGCCTGC GACTTACGGC TCTTTAACAG CGGATAAAAAA AATTATTTG	8820
AAACAAGGTG GATCAGCTAT TTGGCCTTG ATTAGTATT TACTCTTAA ATACACTTGA	8880

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AGGTCGATTC TAATCTCGCT AATCCTTTT AATCCAGAAT AAGGGAAATA TGTTATACTT	8940
GTTCCTTAAGA AAAAAGTCTC ATTGAATTGG TTTTGAGGAG TTAGAAATGA AAGTATTAGT	9000
GACAGGTTTT GAGCCCTTTG GAGGGGAAAA GGCAATCCA CCTTTGGAGG CCATTAAAGG	9060
TTTACCAAGCT GAAATCCATG GTGCTGAGGT CGCGTGGCTA GAGGTGCCGA CAGTTTTCA	9120
CAAATCTGCT CAAGTATTGG AAGAAGAGAT GAATCGTTAT CAACCTGACT TTGTCCTTTG	9180
TATTGGCAA GCTGGTGGAA GAACTAGTTT GACACCTGAA CGAGTGACCA TTAATCAAGA	9240
CGATGCATGC ATTTCTGATA ACGAAGATAA TCAACCGATT GACCGTCCC TTCGCCAGA	9300
TGGTGCCTCG GCCTACTTTA GTAGTTGCC GATTAAAGCG ATGGTTCAAG CTATAAAAAA	9360
AGAGGGCTTA CCGGCCTCTG TTTCCAATAC GGCAGGGACT TTTGTCTGCA GCCATTTGAT	9420
GTATCAGGCT CTCTATTGG TAGAAAAGAA ATCTCCATAT GTTAAGGCAG GTTTTATGCA	9480
TATTCCTTAT ATGATGGAAC AGGTGGTGAA CAGACCGACT ACTCCAGCTA TGAGTTAGT	9540
GGATATTCCG CGAGGGATAG AAGCAGCAAT CGCGCTATA ATAGAACATG GAGATCAGGA	9600
ACTCAAGTTG GTAGGCGGAG AAACTCATTG ATAGAAAAAA GCTTGAGGGG AAAAACCTTC	9660
AAGCTTTGG ACCTTTCCG GCCAATACTG CTCGGTAAA CATAATTAA GTGCATTGGA	9720
TATAAGGTAG GAGTGAAAAA CTAGCAATGC CAAAGGTAAT CCAATTGAGG AAGTACCAAG	9780
GAAGAAGCTG TAAATCTAGG ACAAAAGTGT GGAACCTGTG CCCCTTCATA AAGGAACGGC	9840
TAGTTTTAG GATTCGTCTT GGTGGGACCT GTCCTAGGTC TAGACTATAA CAGACAGAA	9900
ATTCCACCTG TGAATAGGCA TAATACTGTG GAATATAGAG GATATTTCT ACAATGATCA	9960
AGATGAGACT TGCAAGAAAG TAGAGTCCAA AGACCATGAG GAAACGCTCG GTTCAACTG	10020
ATGAGAGATC TAGATTGGA AACTCAGGAT GTAGGGTGAC GAATTTTTG GCTAAAAAGC	10080
TACTATAAAA GAGGAGGTAA ATCCCAAGTA AATTAGGGAT ACTCCATAAA AAGAGATAGA	10140
AACGTTTGAG AACTAGGGTC AAAAAGGTTT GAGAAAAGCG CTCCTCATCA AAGAGAGCTA	10200
GGCTGTTTTT TACAGATGGC TCCGTTTTAG AATCTTCAT GAGTGTCACT GTGCATAGA	10260
CGGAACCTGGT CAAAAGAATA GTCCCGATAA AGGAGACTAG TAGAGGAAAG AGGTAGGTTT	10320
GAAGTATTTG GCCAAGTATG CTGAAAAATG GCTGTTCTAA AACAGTCCCC TGGATCCGAG	10380
ATAAGGGATT AAGAAAACCA GATAAGATGA CCAGCATACT GGGAAGGATA TAGAGGAGAA	10440
AGAGACGGGG GGTGTCAGCC TGAAAATGTT TTGACTCCTG ACGAATTGTT TTTAAATCAA	10500
TTTTTGGATA GTTCATTCTC TTATTATACC ATAGTTCTTA TACATAGTTC GTGACAGTTC	10560
CTACTTTTTT TGATAAAAATC ATACAGTGTG TCCTTGGCA CACTGTATGA ACTGGGACTG	10620

740

TCTTTCCCAG CTTCGGAGGT	AAAAAATGTC AGATTCAACCA	ATCAAATATC	GTTTGATTAA	10680
GAAAGAAAAA CACACAGGGAG	CTCGTCTGGG AGAAATCATC	ACTCCCCACG	GTACCTTCC	10740
GACACCTATG TTTATGCCAG	TTGGGACACA AGCCACTGTC	AAAACTCAGT	CACCTGAAGA	10800
ATTGAAGGAG ATGGGTTCGG	GAATTATCCT ATCAAACACC	TATCATCTCT	GGCTTCGCC	10860
TGGAGATGAA CTCATTGCAC	GCGCTGGTGG TCTCCACAAG	TTCATGAATT	GGGACCAGCC	10920
TATCTTGACA GATAGTGGT	GTTTCAGGT TTATTCTTTA	GCAGATAGCC	GTAATATCAC	10980
AGAAGAAGGA GTAACCTTA	AAAATCATCT AAATGGTCT	AAGATGTTCC	TATCCCCAGA	11040
AAAAGCCATC TCTATTCAAGA	ATAATCTGGG TTCAGACATC	ATGATGTCCT	TTGATGAATG	11100
TCCCTCAGTTT TATCAACCTT	ATGACTACGT TAAGAAATCG	ATCGAGCCGA	CCAGCCGTTG	11160
GGCTGAGCGT GGTTTGAAGG	CTCACCGTCG TCCACATGAC	CAAGGTTTGT	TTGGAATTGT	11220
GCAAGGTGCA GGATTTGAAG	ACCTTCGCCG CCAATCAGCT	CATGATCTG	TCAGCATGGA	11280
TTTCTCAGGC TACTCTATCG	GTGGTTTGGC AGTGGGAGAA	ACCCATGAAG	AGATGAATGC	11340
GGTCTTGGAC TTTACAACTC	AACTGCTGCC TGAAAATAAA	CCTCGTTATC	TGATGGGTGT	11400
GGGAGGCCA GATAGCTTGA	TCGATGGGGT CATTCTGCGG	GTGGATATGT	TTGACTGTGT	11460
CTTACCGACT CGAACATTGCTC	GTAACGGGAC TTGTATGACC	AGTCAAGGAC	GTTCGGTTGT	11520
GAAAATGCC CAGTTTGCTG	AGGACTTTAC GCCACTGGAT	CCTGAGTGTG	ATTGCTACAC	11580
ATGTAATAAC TATACACGCG	CTTACCTTCG TCACCTGCTC	AAGGCTGATG	AAACCTTGG	11640
TATCCGCTTG ACTAGCTACC	ACAATCTTA CTTCTGCTT	AACCTGATGA	AGCAAGTGC	11700
ACAAGCCATC ATGGATGACA	ATCTCTTGGG ATTCCGTGAG	TATTTGTGG	AAAAATATGG	11760
CTATAATAAG TCAGGACGTA	ATTTCTAAAA TGGAATTGAT	ATAAAAAAAT	CCTAAGTTT	11820
CTCTTAGGAT TTTCTTCTT	TTTTTGATAG AATAAAGTGT	ACAATGAAAG	GAAGAATAAA	11880
CTCGTATGCG CATTAATGG	TTTCCTCGA TTAGG			11915

(2) INFORMATION FOR SEQ ID NO: 97:

- (i) SEQUENCE CHARACTERISTICS:
- (A) LENGTH: 9069 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 97:

GAGAGGGCAA CAGTTCTATC	GCTTCAAATT TTTTCTTGGT	TTGCAGATAT	TCAAGAACG	60
GGAGTTTTTC TATAGTATTC	GGCAGATTAA TTACAGCCAA	GCATCTCAA	AATACGGACA	120

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GCATCCTCCA	TCTTTTCTG	GCCTTCCTTG	ACTCTACCTT	GCTTGCTATC	AAGGAGACCT	180
TCTGCCACA	GATAAACAAAT	TCGGAAATAG	GTCTCATTTT	CCTTGTAGAA	ATGCTCTTCG	240
ATAACACGTT	TAAAATAATA	GGCATTGGTA	AATTCTTCAC	ACTCAATACT	AGCTAAAAAG	300
CCATTCAATA	GTATAGTATG	AAAAAGGTTT	CGATTGCCAG	ACATTTCCAT	TAGAAAATCA	360
GATTACGTA	CCATTCTCG	TACATATCTA	GTAAAAAGAG	AAACAGATAA	AAATGGAGAA	420
CTGACTGAAA	ATAAATTGAG	TTCATAGATT	CCCCAGATCT	CGGTAGAAAA	CAAATAATCA	480
TGAAGGACTT	TTCCCTCCTC	TGCTGTTAAG	TCTACCCTTT	CATCTATGCT	CTTCATATAA	540
GACTGATAA	TAATGGCATT	TAGAATATGT	TTCTGTTGT	TGTGAGAAATG	GGCATGCTTT	600
TATACTCCCT	GCGATATAAG	TCCTCAAGAG	GTGCTATATT	CTTGTTTCC	AAGACATCTG	660
TAATTCTTT	TCTCAACTCA	GAATCTGTAT	CATACTGGAA	ACCTCTGCC	AGAAAAGAGGA	720
TCTCCTCCAC	ACTGGCAGAT	ATATTTCCA	GAGCAAATAG	AAACTTTCC	ACCGAAAGCT	780
CACTCTGACC	TGTTTCAAAA	CGGGACAACA	TAGACGGCGA	AAATTGTCCT	CCGGTTGCTT	840
GTCTCAGTGA	GATATTCTT	GACTCTCGTA	ATTGTCTAAA	GACTTTCCA	ATCTGCTCCA	900
TAGACTTCCC	CTTGATTCCG	TATTTCTTC	ATTTTATCAT	ATTTTCAGA	AAATTCTATCA	960
AAAACTTGCC	AAATTGTCAG	AATTATGAGA	AAATAGAGGA	TATTTATCAC	GTGGAGGGAC	1020
TGCTATGAGA	GACGATATCA	AAATCAATGA	CCGTGCTTTG	GCCTGCAAG	ACCAAATTAT	1080
CGAAAAAACTA	GAGAAAGTTT	TTGATACAGA	TGTGAAATTG	GATCTTACA	ATCTAGGTCT	1140
GATTATGAA	ATCAATCTGG	ATGAAACGGG	GCTCTGCAAG	ATTCTCATGA	CCTTCACCGA	1200
TACTGCCTGT	GATTGCGCCG	AAAGCCTGCC	TATTGAAATC	GTGGCAGGTC	TGAAACAAAT	1260
CGAGGGTATC	AAAGATATCA	AGGTTGAAGT	TACCTGGTCG	CCTGCTTGGA	AAATCACACG	1320
AATCAGTCGC	TATGGCCGTA	TTGCCCTTGG	ACTACCACCT	CGTTAAGCAG	ACCAATCACT	1380
TTTAAAGATG	AAAATCAAAG	GCAAACTAG	AAAACTAGCC	GCAGGTTGCT	CAAACACTG	1440
TTTTGAAAGTT	ATGGATAGAA	CTGACGAAGT	CAGCTCAAAA	CACTGTTTG	AGGTTGTGGA	1500
TAGAACTGAC	GAAGTCAGCT	CAAACACTG	TTTGAGGTT	GTGGATAGAA	CTGACGAAGT	1560
CAGCCCCAAA	CACTGTTTG	AGGTTGTGGA	TAGAACTGAC	GAAGTCAGTA	ACCATAACCTA	1620
CGGCAAGGCG	ACGTTGACGT	GATTTGAAGA	GATTTTCGAG	TATGAGTTA	TTTTTACCT	1680
GACTTGTCCA	TATTCCAGAA	GTCTGTCACG	GCTCCGCGTG	AAGCAGATGA	TACGATGTGG	1740
GCATATTTCAC	CGAGGGACACC	ACGGCTGTAA	AGTGGTGGCA	AGGTTGTTTC	TGCCTTGCGT	1800
TTTTCAAGTT	CTTCTTCGGA	TACGGCCATA	AAAATTCTT	TGGTATCTTG	GTCAACCGTA	1860

742	
ACGATATCGC CGGTACGGAG ATAGGCAATT GGTCCACCAT CCTGAGCTTC AGGAGCGATA	1920
TGTCCAACAA CCAGACCATA AGTACCACCA GAGAAACGTC CGTCCGTCAA GAGGCCACC	1980
TTATCTCCCT GACCTTTACC ACAATCATT GAAGAAAGTG ATAGCATCTC AGGCATACCA	2040
GGACCACCTT TAGGTCCAAC AAAACGAACA ACGACTACAT CGCCATCAAC GATTCATCT	2100
GTCAGAACGG CCTGAATCGC ATCTTCTTCT GAGTCAAAGA CCTTAGCTGG CCCAACGTGA	2160
CGACGCACTT TAACACCTGA TACCTTGGCA ACTGCACCGT CAGGAGCAAG GTTCCCCTTC	2220
AAGATGATAA CGGGACCATC CGCACGTTT GGATTTCAA GTGGCATGAT AACTTTTGG	2280
CCTGGAGTCA AGTCTGCAA GTCAGCCAAG TTTTCAGCTA CAGTCTTAC AGTACATGTG	2340
ATGGATCTC CGTGAAGGAA ACCATTTGCC ACAAAATACT TCATAACCGC AGGGACACCA	2400
CCGACTTCGT AGAGGTCTTG GAAGACATAC TGACCAGATG GTTCAAGTC GGCAAGTGA	2460
GGCACACGTT CTTGAATCGT ATTGAAGTCC TCAAGTGACA AGTCAACATT TGCGGCATGG	2520
GCAATGGCGA GCAAGTGAAG AGTGGCGTTT GTAGAACAC CGAGAGCCAT CGTTACAGTG	2580
ATAGCATCTT CAAAGGCTTC ACGAGTCAAG ATATCTGATG GTTGAGACC AAGTTCCAAC	2640
ATCTTAACAA CAGCACGTCC TGCTGCTTCG ATATCTTCTT TCTTATCAGC TGATTCAAGCT	2700
GGGTGAGAGG ATGACCCCTGG CAAACTCATC CCTAGAACTT CGATAGCAGT TGCCATGGTA	2760
TTAGCAGTAT ACATACCAACC ACAACCACCA GGGCCAGGGC AGGCATTACA TTCAAGACGT	2820
TTCACGTCTT CAGCTGTAT GTCACCGTGG TTCCATTTC CGATACCTTC AAAGACAGAA	2880
ACCAAGTCGA TATCTTACCA ATCAAGATTT CCCGGTGCAA TAGTCCACC ATAGGCAGAA	2940
ATAGCTGGGA TATCCATATT AGCAATAGCA ATCATAGATC CAGGCATGTT CTTGTCACAG	3000
CCACCGATAG CGACGAAGGC ATCCACGTT TGACCACTCA TAGCCGCTC GATGGAGTCC	3060
GCGATGATGT CACGAGATGT TAGAGAGAAA CGCATACCAAG GCGTCCCAT AGCGATCCCG	3120
TCCGCTACGG TAATGGTTCC AAACTGTACA GGCAAGCGC CTGCAGATTT GACACCTTCT	3180
TTAGCCAGTT TCCCGAAATC ATGCAAGTGA ATGTTACATG GTGTATTTTC CGCCAAGTC	3240
GAAATCACTC CCACAATCGA TGTTCAAAG TCCTTATCTG TCATACCAAGT CGCACGAAGC	3300
ATAGCACGGT TAGGTGATTT ACCATGCTG TCATAATGC TACTGCGGTG ACGTTTATCT	3360
AATTCACTA TCTTATCCCT CCCATTTCAAG TTTTTACTAT TATAGCACAA TTTTCGCATG	3420
AAGAACAGAA TAAAATTCTT GAATTTTCAG AAAATTCTAT ACACATGTGA AATATTTAAA	3480
ATTAAAAACA ACAAAGCGGA TTAGTGCAC TCTGATGAC CAGAATATGC TTTTTAATCC	3540
GCTTTCTTTA AATAACGTAC TGTATTTTT ACAGAAATTC TTTCAAATAA GTGTATTTAA	3600
CATCTATCTT GCATTATAAA TTTCTAGAAC CTTCTCTTTT ATATTCGATT CACTCAAAAC	3660

743

ATACTCATTA AGAAGATAAT CCATTTCCC TACTTGACCG AATCTTCTT GAACACCCAT	3720
CCGATGAATT TTTGTTATTC CATCATCAGA GAATAATTCA CATAAAGCAC TGCCAATTCC	3780
ACCTATCTGA TTGTGGTTT CTACAGTAAA TATAGTTTT CCACTTAACAA TTGTTTTAT	3840
CTGTTCTGGT ATCGGTTGA TTCTAAATAA ATCTATCACAA CCTACTGAAT AACCTAATT	3900
AGACAGTTCA TCTGCAACTC GAATACTTGG AGCAACCATT ATGCCAGAAG CAACGATTAC	3960
AAGATCTTCA CCATGCCTTA ACTCAATGTA GCCTTTAGAA AAATCTTCTC CACCTTGATA	4020
CACAGGAACT GGAGCTTTTC TAATTGTTCG AATATATTTT AGTCCTTTA AGTCTAATGT	4080
CTGGTTCAAT ATTTCACGAA ATTGGATATC ATCAGTTGCT TCGAAATGA TTGATTTAGG	4140
AATTAACGT ACAATCCAA TTTCTTCAAA TGGCATATGT GTTCCACCAT TCATCTCTGC	4200
CGTTACTCCT GCATCTGATC CAATCACAGT GGCATCCAAT TGTGCGTATC CAAGAGAAAT	4260
AAATAATTGA TCAAATACTC TTCTGTGAAAG AAAAGGACCA AATGTATGAA GATAAGGTCT	4320
AAACCCCTGA ATAGACAAGC CTGCTGCAAG GCCGACCATT TCTGCTTCCA TAATCCAAAC	4380
ATTCACATAA CGGTCTCCAA AGTCCTTTTC AAGATTATTA GTAGCCATCG AACTTGACAA	4440
ATCGGCTTCT AAGACTACTA TATCAGAATC ACTTTGATTA GCCTCTAAAA GGAAGTCTCT	4500
ATATACATGC CGTAATTCTT TCGTACTTCT CATCATTCTG TTTCTCTCAA TTCCTGACTT	4560
AATCTTCTA CAACTGAAGT TAACATTGT TTCTCTCTA CAGTAGGGCG AAGATGATGA	4620
TTGGATTTCAG TTTCTCCAG CTCTGTGAAACC CCTTGACCTT TAATAGTATC TAATACAATG	4680
CACTTAGGTG ATGAAATTATT TGACTGTTT AATGGACAA TCCCTTCATA AATTCTCTA	4740
ATATCTGAAC CCTTGACCCCT AATGGATTCA AATCCAAATG CTGAAATTT TTCTACGAAA	4800
TCACCTGGAT TACAATATC CTTTGTAAAA CCATCTAATT GTTTTTGTT ATCATCAACA	4860
AATACAATTA AGTGGATAA CTGTTGATGA GAAGCAAACGT GTATAGCCTC CCAACATTGT	4920
CCCTCATTTA ACTCACCATC TCCAACAATA GCGTAAGTAT AAAAGGACT CTTCTTATT	4980
CTCTGACCAT ATGCAAGTCC AGTTGCAACA CTAATTCTT GTCCTAAAGA GCCCGTTGTC	5040
ATATCTATGC CTGGCGTTAG ATTTCTATCA GGATGAGACCG GTAATTTGGT TCCATTGTA	5100
TTTAAAGAAT ATAAGAATTC TTGTCAAAG AAACCATTCA AATAGAGTGT ACTGTATAGA	5160
GCTGGTCCTC CGTGACCTTT TGATAATATG AAATAATCTC TATCTCGTGC TGCAAATATT	5220
TCTGGAGTCA TTGGCATTAT TTCAACCTAA AGCACCGCTA AAACCTCTAC GATAGACAGA	5280
CTTCCTCCGT AATGTCCGAA TCCAAGATGA TTCAATGTMTC TAAGAGTATT TAATCGGATG	5340
TTAGTCGCAA ATTTCTTAA CCCATCTTCT CTATTTTAC TAAATCAT CCCTTATTCC	5400

744	
TCCGTTGCAG ATGGCTTTT AATAAAGGAT ACTCCAACA TAACTGCTAG AATAAGAAC	5460
AGACCAATCA CAATGCCGTGC TTGTGAGCCA AATTGATTAA ACATTCCTAA AATAATTCC	5520
GATAGACCAA AATCTGCATC TGAGAAAGTT GATCCTTGGAA ACCAACGTCC TCCCAAA	5580
GGCATTAAAA AGACTGGAAG AAAACTGATT AAAATACCTT GTAAAAATGC TCCAATAGTG	5640
GCTCCACGAA CACCACAGA TGCATTCCCA ATGACACCTG CAGTCGCTCC ACAGAAGAAA	5700
TGAGGCACAA CACCTGGTAA GATAACAACC GTTCCGTAAAG CAATCATATAAT TACCATACTT	5760
ACTAAACCAC CAACAAAAGT AGAGATAAAAT CCAATTAGAA CTGCATTGGG TGCATAAGTA	5820
TAAACAATCG GACAATCCAA AGCAGGTTTT GAATTAGGTA CAAGACGCTC TGAAATACCT	5880
TTAAAGGCTG GAACAATTTC GCCAAAATA AGGGAACAC CTGCTAAAAT AACAAATACC	5940
CCTGCTGCAA ATTGACCTGC TAATTGTAAA GCATAAACTA GACCACTTGT ACCACTACTG	6000
ATTTCTTTTT CTATATATTC TGACCCCTGCA AAGATAGCTA CAATAATGTA AATAACTGCC	6060
ATGGATAAAAG TAATACAACT AGTACTATCA CGTAAAAAAAG CTAAACTCTT TGGAAATTAA	6120
ATGTCCTCTG TTGATTTGA TTTGTCACCG ATAAGGCTAC CAGTAAAACC ACTCAACCAA	6180
TATCCCCAAAG AACTGAAATG ACCTAAAGCT ACCTTGTCA TTCCAGTTAA TTGAACCATA	6240
TATTTTGCA CAAATGCTGG GGAAATACTC ATAATAATAC CGAGTGCTAA TCCTCCTAGT	6300
AAGATGAGAG GCAAGCTACT AAAGCCAGCA ACTGATAAAA TGACCGCAAT CATACTGCC	6360
ATATATAGAG TGTGGTGCCTC TGTTAAAAAA ATATATTAA ATCGAGTAAA ACGACCGATT	6420
AAGATATTGA ACACCATGCC TGCAAAACATA ATCATTGCAG TAGCTGAGCC ATATGTTGTT	6480
AAAGCTACAG CTACAATTGC TTCATTATTC GGACACACGC CAGATAATG AAAAGCATGC	6540
TCAAACATGG TACCAAAATGG ATTTAAAGAA TTTTGTACAA TTCTGCACC ACCAGATACA	6600
ACTAAGAAAC CAACAAAGGT CTTAATTCCA CCTTTAATAA TATCAGGTAA TTTCTCTTC	6660
TGAAGAACTA ATCTTAAGAT TGCAATTAAA GCTACTAAAA TAGCTGGTGT ACTAACAAATA	6720
TCCAATATGA ACTTCATCAT GACGCTAGCC TCCTATATAA GTCTTTTTC TTCAACAAAGT	6780
TTAGTAATTA ATTCTCGTAG TTTCATCCATA TCAATAATAC TATTTAAGAT ACGAACATCT	6840
CCAAGATGAC TAGCTGAATC AGCTAGATCA CGACCAACAA TCCAAATATC AGCTGCATT	6900
GGATCTGCTC CACCTAAATC ATAATGTTCA ACTTCTACAT CCGAAACATT CAAATCACTC	6960
AATACAGATT CAATATTTCAT CTGTACCATATAA AACTTGAAAC CTAATCCTGA ACCACAAAGCT	7020
GTACCAATT TTAACATTAT CTAATCCTCC TGTTTAATTA TCATTTTAAT GTCATCATAG	7080
TTTTTGATG ATATTAAGT TTGAACATGA TTTTTATCTC TTAAAATTGT TGTTAAATGT	7140
GACAAAGCCT TAAATGACT CTCATTATCA ATGGCTGCAA TACAAATCAA CAATCTTACC	7200

745

TCTTGTTCCTG GATTATCCAA TAAATAAATC GGTTCTTCCA AACTAACAT TGACATTCCT	7260
ATTCATTCA CACCTTCATC TGGCCGAGCG TGAGGAATTG CTACTCCCTT CCCTAAATTA	7320
ATAAAAGGTC CAAACTCTTC TACTTTTGAT ATCATTGCCCT CAGGGTAGTT CTCAGTTATC	7380
TTATCTTGAT CAAAAGCGG TTTAGCTGCT AAACGAATCG CCTCCCTTCCA TCCTAATTTT	7440
TGCGAACTAA CCTGATAGGT TTCTTTGGTA ATAAGTTGTT CTAGCACTGG TACAATTCC	7500
TTTCTATCAT TTTTTGGTA AAGATAATTC TTTAACGCCA ATCTTAATTC CAATTCTTGT	7560
GTAATAATTC CATATCTTTT GACAATATTC AGGATTGTT CAATCTCAA ATCTCCATAC	7620
TCTAAATTGTC GAAAATCTTT TAACACTAGT TCTACTAGTT GTATTGCTTG CTCTTCAGTC	7680
ATCATAACCG AAACTAGATA ATTTGGCTTT TCTGTCTCCA CCTTTATGGT AGAAAAAACCC	7740
ATATCATAGT CACTACTAGC TTTCACCTGT AAATCATCAA TCTTTGAGGT TCCTATAAAC	7800
TCAATTGAG GAAATAATGC TAATAGATTC TCTTTAACCA TCAATGAAGA ACTAACACCA	7860
TTAGGACAAA TGATTGCTGC TTTATACCAT TTTTGAGGCA AAGTATCTGC TTTCTTTAAA	7920
TAACCTCCGA AATGGATAAC AAAATATGCT GTTTCACTAT CAGGTATGGG ATTGTCAATA	7980
GCCTCCATCA AGGGCATCAA AGAATCTTG ACTAATTCAA ATAATCAGG ATAATGTTCT	8040
TTAACATGCA ATACATATTC ATTTGAACTA GGTAGGCCGA ACTTTAATCT ATAGTAAGCC	8100
GGTATAAGGT GCGGGCGAAG ATTTCTCTC AATCCTTCCC TTTGTTAAA ATGTAACAAA	8160
GAAATATCTT CCATTCTACT TATAATAGCC TCTGTTAATT GATTAAAGTA AACCCCAACCA	8220
ACATCTACTT CACCTTCAAA GCAACTTGAT AATAAAACGG TGATATAGCG ATAATCATCC	8280
TCAGAAAACA CCGTATCTAT AATTCCCCAA TCAACCACTG TATCCAATTA AATAGTGGTT	8340
ATATCTTGAA TAACAGGAGA TACTAATGTC TCTGAAAGAC ATACTCTTTC AACATCCCTT	8400
TGATACCTAC ACAGAATGAA TACTAAACCG AAAAGTAAA CTTTTAATTG ATTAACAAATA	8460
GGTACTAGCT GTAGCTCTC ATAATAATCT TTAACACTT GATCAATCAA ATCATAAGTT	8520
AATGAATACC CCCAACTGGA TAAAACATAA TCCAAACCCC AAATCCCTAT GGAGGATTCC	8580
AGCAACTCAC TAACCATTG AAAAGCTAAG CGGTGCTTAT TCCACTCTGA ACCGTGTAAA	8640
GTATAACCTT TTGCTCTACT GTACCCTAGC TCCAAATCAT TATCTAACAT AATCTTCTT	8700
AATGATTGAA TATCAGATAA GGTTGTATTC TTACTTACTT TCAAAAAGTC TTGGTAATGA	8760
CTATTGATA TAAAATCTAA TCGGCAAAAA GTGTAAAGAT AGATTAAGC TAAGCGAGTC	8820
GACTTTGGTA AAACCAATTC ATCCGACTTA ATAATATCTG TCAAAGACTG CTTCGTACGA	8880
TTTGATAAAC TATAGCGACC TTGCTTTTA TCCAGCACTA TCCCTTTATT AGCTAGATAA	8940

746	
GGCACTAAAT AATCTATTCC TTCTTTGACT TCCTTATAG GTAAGCTCAC CTTAACAGAT	9000
AATTCATATA ACGATAGCTC ACAATGATCC ATCAAAGTCA TCAAAATAAC TAGTGCTCTA	9060
TAATCAAAC	9069

(2) INFORMATION FOR SEQ ID NO: 98:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 8654 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 98:

CGAGACAACA AGATGAAGAA AAATTTGCC TATCGTTCTG GGCCTTGCA AGTGTAGCAC	60
TTCTTGCAGC CTGTGGAGAA GTGAAGTCTG GAGCAGTCAA CACTGCTGGT AACTCAGTAG	120
AGGAAAAGAC ATTAAAATC GGTTTAACT TTGAAGAAC AGGTTCTTA GCTGCATACG	180
GAACAGCTGA ACAAAAAGGT GCCCAATTGG CTGTTGATGA AATCAATGCC GCAGTGGTAT	240
CGATGGAAAA CAAATCGAAG TAGTCGATAA AGATAATAAG TCTGAAACAG CTGAGGCTGC	300
TTCAGTTACA ACTAACCTTG TAACCCAATC TAAAGTATCA GCAGTCGTAG GACCTGCGAC	360
ATCTGGTGCG ACTGCAGCTG CGGTAGCGAA CGCTACAAAA GCAGGTGTTG CATTGATCTC	420
ACCAAGTGCG ACTCAAGATG GATTGACTAA AGGTCAAGAT TACCTCTTA TTGAACTTT	480
CCAAGATAGC TTCCAAGGAA AAATTATCTC AACTATGTT TCTGAAAAAT TAAATGCTAA	540
GAAAGTTGTT CTTTACACTG ACAATGCCAG TGACTATGCT AAAGGGATTG CAAAATCTTT	600
CCGGAGTCA TACAAGGGTG AAATCGTTGC AGATGAAACT TTCGTAGCAG GTGACACAGA	660
CTTCCAAGCA GCCCTTACAA AAATGAAAGG GAAAGACTTT GATGCTATCG TTGTTCTGG	720
TTACTATAAT GAGGCTGGTA AAATTGTTAA CCAAGCGCGT GGCATGGGAA TTGACAAACC	780
AATCGTTGGT GGTGATGGAT TCAACGGTGA GGAGTTGTA CAACAAGCAA CTGCTGAAAA	840
AGCATCAAAC ATCTACTTTA TCTCAGGCTT CTCAACTACT GTAGAAGTTT CAGCTAAAGC	900
TAAAGCCTTC CTTGACGCTT ACCGTGCTAA GTACAATGAA GAGCCTCAA CATTGCGAC	960
CTTGGCTTAT GATTCAGTTC ACCTTGTAGC AAACCGCAGCA AAAGGTGCTA AAAATTCAAGG	1020
TGAAATCAAG AATAACCTTG CTAAAACAAA AGATTTGAA GGTGTAACGT GTCAAACAAG	1080
CTTCGATGCA GACCACAAACA CAGTCAAAAC TGCTTACATG ATGACCATGA ACAATGGTAA	1140
AGTTGAAGCA GCAGAAGTTG TAAAACCATA ATAGAAAAAT GTTGAAATAG GGAATGAGCC	1200
TTTGACTCAC TCCCTGTTTC GATTTTAAT ACTCTTCGAA AATCTCTCA AACTGCGTCA	1260

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ACGTCGCCCTT GGATTATATA TGTGACTGAC TTCGTCAGTC TTATCTACAA CCTCAAAGCA	1320
GTCCTTGAG CAACCTGCGG CTAGTTCT AGTTTGCTCT TTGATTTCA TTGAGTATAA	1380
GAACCTATCA AAAAGTGAGG GAAAACCCTC GGAATTATAA ATAGAAAGAG TGAATCTTAT	1440
GCTCCAACAA CTCGTAATG GTTTGATTCT AGGTAGTGT TACGCGCTGT TAGCCCTAGG	1500
ATATACCATG GTTTACGGAA TTATCAAGCT CATCAACTTC GCCCATGGTG ATATTTATAT	1560
GATGGGAGCC TTTATCGGTT ATTTCTTGAT CAATTCTTTC CAAATGAATT TCTTTGAGC	1620
GCTTATTGTA GCTATGCTAG CGACAGCTAT TCTTGGTGTG TGATTGAGT TTCTTGCTTA	1680
CCGACCTTTG CGCCACTCTA CTCGTATTGC TGTTTGATT ACGGCTATTG GGGTTCTTT	1740
CCTATTGGAG TATGGAATGG TCTATCTGGT TGGTGCCTAAT ACCCGTGCCT TCCCCTAACG	1800
GATTCAAACA GTTCGATATG ATTTGGGACC AATTAGCTTA ACAAATGTGC AGTTAATGAT	1860
TTTGGCCATT TCCTTGATTT TGATGATTTT GTTACAAGTC ATTGTCCAAA AGACTAAGAT	1920
GGGGAAAGCC ATGCGTGCAG TATCAGTAGA TAGCGACCGG GCGCAATTGA TGGGGATCAA	1980
TGTAACCCTG ACGATTAGCT TTACCTTCGC TTTGGGTTCT GCTCTTGCCT GTGCCGCTGG	2040
TGTTCTGATT GCTCTTTATT ATAACCTCTCT TGAGCCTTTG ATGGGGTTA CTCCAGGTCT	2100
TAAATCTTC GTTGCCCGCAG TACTTGGTGG TATCGGAATT ATTCTGGTG CGGCTCTTGG	2160
TGGCTTG TG ATTGGTCTAT TGGAAACCTT TCGGACTGCC TTTGGGATGT CAGATTCCG	2220
TGATGCCATT GTTTATGGAA TCTTGGTGT GATCTTGATT GTCCGCCAG CTGGTATCCT	2280
TGGTAAGAAT GTGAAAGAGA AGGTGTAAAC GATGAAGGAA AATTAAAAG TTAATATTCT	2340
ATGGTTACTC CTTTTGGTAG CTGGCTATAG CTTGATTAGT GTACTGGTTT CAGTCGGAGT	2400
ACTTAATCTA TTCTATGTAC AGATTTACA ACAAATTGGA ATTAATATTA TTTTGGCTGT	2460
TGGCTCAAC TTAATCGTG GTTTTCAGG ACAATTCTA CTTGGTCATG CTGGTTTCAT	2520
GGCGATTGGT GCCTATGCAG CAGCTATTAT TGGTTCTAAA TCACCAACCT ACGGTGCCCT	2580
CTTTGGAGCT ATGCTTGAG GGGCTTGCT TTCAGGAGCA GTTGCCTTAC TTGTCGGCAT	2640
TCCAACCTTG CGCTTGAAGG GGGACTATCT TGCGGTAGCA ACTCTGGGTG TTTCTGAAAT	2700
TATCCGTATC TTTATCATCA ATGGTGAAG CCTTACAAAT GGTGCAGCAG GTATCTTAGG	2760
GATTCCTAAC TTACAACCTT GGCAAATGGT TTACTTCTTT GTCGTGATTA CAACCATTGC	2820
AACCTTGAAAC TTCTTGCCTA GCCCAATTGG TCGTTCAACC CTCTCTGTTC GTGAAGATGA	2880
AATCGCTGCT GAGTCAGTTG GGGTTAATAC GACTAAAATT AAAATCATCG CTTTTGTCTT	2940
TGGTGCCTT ACTGCAAGTA TTGCTGGTC ACTTCAGGCA GGATTTATCG GGTCTGTTGT	3000

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ACCGAAAGAT TACACCTTCA TCAACTCAAT CAACGTTTG ATTATTGTTG TATTTGGTGG	3060
ACTCGGTTCC ATTACAGGTG CGATTGTTTC GGCTATTGTT CTGGGAATT TGAAATATGCT	3120
TCTCCAAGAT GTTGCTAGTG TGCCTATGAT TATTTACGCT TTGGCCTTGG TATTGGTAAT	3180
GATTTTCAGA CCAGGTGGAC TCCTTGGAAC ATGGGAACCT AGCCTATCAC GTTTCTTAA	3240
AAAATCTAAG AAGGAGGAAC AAAACTAATG GCATTACTTG AAGTAAAACA GTTAACCAAA	3300
CATTTGGTG CTCTAACAGC TGTTGGAGAT GTGACTCTTG AATTGAACGA AGGGAACTG	3360
GTTGGATTAA TCGGTCCAAA CGGAGCTGGG AAAACCACCC TTTCAACCT TTTGACCGGT	3420
GTTCATGAAC CAAGCGAGGG AACAGTAACC CTAGATGGTC ACCTTTGAA TGGGAAATCA	3480
CCTTATAAGA TTGCCTCTTT GGGACTTGGGA CGTACTTTCC AAAATATCCG TCTCTTTAAA	3540
GATTTAACAG TTTTAGATAA TGTTTGATT GCTTTGGAA ACCATCACAA ACAGCATGTT	3600
TTTACTAGTT TCTTACGCTT ACCAGCTTT TACAAGAGTG AAAAAGAATT AAAGGCTAAA	3660
GCTTTGGAAT TGTTGAAAAT CTTTGATTAA GATGGTGTG CAGAGACTCT TGCTAAAAAT	3720
CTTCCCTACG GACAACAACG TCGTTTGAA ATTGTTCGTG CCTTGCTAC GGAACCTAAA	3780
ATTCTCTCT TAGATGAACC AGCAGCAGGT ATGAACCCAC AGGAAACAGC CGAATTGACT	3840
GAGTTAATTC GTCGTATCAA AGATGAGTTT AAGATTACAA TCATGTTGAT TGAACACGAT	3900
ATGAATCTGG TCATGGAAGT AACAGAACGT ATCTACGTAC TTGAATATGG CGTTTAAATC	3960
GCTCAAGGAA CTCCAGACGA AATTAAGACC AATAAACCGC TTATCGAAC TTATCTAGGA	4020
GGTGAAGCCT AATGTCTATG TTAAAAGTTG AAAATCTTC TGTGCATTAC GGTATGATCC	4080
AAGCAGTTCG TGATGTAAGC TTTGAAGTTA ATGAAGGAGA AGTTGTTCC CTTATCGGTG	4140
CCAACGGTGC AGGTAAGACA ACTATTCTTC GCACCTTGTC AGGTTGGTT CGACCAAGTT	4200
CAGGAAAGAT TGAATTTTA GGTCAAGAAA TCCAAAAAAT GCCAGCTCAG AAAATCGTGG	4260
CAAGTGGTCT TTCACAAGTT CCAGAAGGAC GCCACGTCTT CCTGGCTTG ACTGTTATGG	4320
AAAATCTTGA AATGGGAGCT TTCTTAAAGA AAAATCGTGA AGAAAATCAA GCTAACTTGA	4380
AGAAGGTTTT CTCACGCTTT CCTCGTCTTG AAGAACGGAA GAACCAAGAT GCAGGCCACTC	4440
TTTCAGGGGG GGAACAAACAA ATGCTTGCCA TGGGACGGCG CCTCATGTCA ACACCAAAAC	4500
TTCTCTTTT AGATGAACCA TCAATGGAC TTGCCCCAAT CTTTATCCAA GAAATTTTG	4560
ATATCATTCA AGATATTCAAG AAGCAAGGAA CAACGGTCCT CTTGATTGAA CAAAATGCCA	4620
ATAAAGCACT TGCAATCTCT GACCGAGGAT ATGTAATGGA AACAGGGAGA ATCGTCTTAT	4680
CAGGAACAGG AAAAGAACTC GCTTCATCAG AAGAAGTCAG AAAAGCATAT CTAGGTGGCT	4740
AAAACAATCC AGTGGATTGT TTTAGTCGGC AGATGGAGAT TACGAAGTAA TCATCAATAT	4800

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AGTCCGGGGG ACCTTTTAG TCGGTAGATT GAGATTGCAA ACAAATCTGC ATCTACATTG	4860
AAAGCTTAAT TTCTAATAAT TGAAAAAATC GAATGAAAAA TTTCTTACCT TCATTACAG	4920
AGCTCGATTT CAGAGCTCTT TTTGCTAGCT TATTCATACT TTTCTGAATT TCGAAAAGA	4980
AATGTAAGCG TTTGATAGAT TTACAAAAG ATTGTATAAT AGGGATAAGA ATAGAAAAGG	5040
AGAAGTCTCA TGGCAGTTAA AGATTTATG ACCCGCAAGG TAGTTTATAT TAGTCCAGAT	5100
ATAACAGTAT CTCATGCAGC AGATTGATG AGAGAGCAAG GTTGCACCG TCTGCCTGTT	5160
ATCGAAAATG ATCAATTAGT TGGTTGGTG ACTGAGGGAA CCATTGCACA AGCAAGTCCA	5220
TCTAAAGCAA CAAGTCTTC TATCTATGAG ATGAATTATC TTCTGAATAA GACAAAAGTA	5280
AAAGATGTCA TGATTGCGA TGTTGTCACT GTCTCAGGCT ATGCTAGTCT AGAAGATGCA	5340
ACTTATCTGA TGTGAAAAA TAAGATTAGT ATTCTCCCTG TCGTAGATAA CCATCAAGTA	5400
TACGGAGTTA TTACTGACCG TGACGTTTC CAAGCCTTTC TTGAAATTGC AGGTTATGGC	5460
GAAGAAGGGA TTCGTGTACG CTTTGTACCA GAAGATGAAG TTGGTGTCT TGGAAAATT	5520
GTTTCTTGA TTGAGAAGA AAATTTGAAT ATCTCCCATA CAGTCAATAT TCCGCGTAAG	5580
GATGCTAAGG TGATTATCGA AGTGCAAATC GATGGATCAA TTGATTTACC AGCCTTGAAA	5640
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TTGTAAGAAG GGAAGCCAA AGGCTCTTT TTTCATGAA AGGGGATTAG AGCAAAAGAT	5760
GGAAAGAAAT GATAAAATAT GCTATAATGA AATAATGAA AAAAGGAGTA TTTATGGACA	5820
TTTCAGTAAT TCGTCAGAAA ATTGACGCAA ATCGTAAAA ATTAGCTTCT TTCAGGGGT	5880
CTCTTGACC TCGAAGGGCT AGAGGAAGAG ATTGCCATCT TGGAAAACAA GATGACAGAA	5940
CCTGATTTT GGAACGATAA TATTGCGGCC CAAAAAACGT CGCAAGAATT AAATGAATTA	6000
AAAAACACTT ACAATACCTT CCATAAGATG GAAGAGTTGC AGGATGAAGT CGAAATTTA	6060
TTGGATTTT TGGTGAAGA CGAGTCAGTG CATGATGAAC TGGTAGCGCA GTTAGCCGAA	6120
CTTGATAAGA TAATGACCA CTACGAGATG ACTCTACTCT TGTCAGAACC TTATGACCAC	6180
AACAATGCCA TCTTGGAAAT CCATCCAGGT TCTGGTGGTA CTGAGGCGCA GGACTGGGT	6240
GATATGTTGC TTCCTATGTA TACTCGTTAT GGTAATGCTA AAGGCTTAA AGTGGAAAGTG	6300
TTGGATTACC AAGCAGGTGA TGAGGCTGGT ATTAAGTCGG TAACTTTATC ATTTGAAGGG	6360
CCTAATGCCT ATGGTCTCCT CAAGTCAGAA ATGGGTGTC ACCGCTTAGT GCGAATCTCA	6420
CCATTGACT CTGCCAACG TCGCCATACC TCTTCACAT CTGTAGAAGT GATGCCAGAA	6480
TTGGATGATA CTATTGAAGT GGAAATCCGT GAAGATGATA TCAAGATGGA TACCTCCGT	6540

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TCAGGTGGTG	CCGGTGGACA	AAACGTCAT	AAGGTTCAA	CAGGTGTACG	TTTAACCCAC	6600
ATTCACAATG	GAATTGTTGT	CCAATCAACA	GTAGATCGTA	CCCAGTATGG	AAATAGAGAT	6660
CGTGCATGA	AGATGTTGCA	GGCTAAGCTC	TATCAAATGG	AGCAAGATAA	GAAGGCTGCG	6720
GAGGTAGATT	CTCTCAAAGG	TGAGAAAAAG	GAGATCACTT	GGGAAAGCCA	AATCCGTTCT	6780
TATGTCTTCA	CCCTTATAC	TATGGTAAAA	GATCACCGAA	CTAGCTTGA	GGTTGCTCAG	6840
GTAGATAAGG	TTATGGATGG	GGACCTAGAT	GGTTTATCG	ATGCTTATCT	CAAGTGGCGA	6900
ATTAGCTAAG	ATAGAAAGGA	ACTCACATGT	CAATTATTGA	AATGAGAGAT	GTCGTTAAAA	6960
AATACGACAA	CGGAACAAC	GCTCTACGCG	GTGTTTCGGT	TAGCGTTCAA	CCGGGGAAAT	7020
TTGCTTACAT	CGTAGGACCT	TCAGGAGCAG	GGAAAGTCAC	TTTTATTCTG	TCTCTGTATC	7080
GTGAAGTAAA	AATCGATAAA	GGAAGCCTAT	CAGTTGCTGG	TTTTAATCTG	GTAAAGATCA	7140
AAAAGAAAGA	TGTCCCGCTT	CTACGTCGTA	GTGTTGGGGT	TGTCTTCCAG	GATTATAAAT	7200
TGTTACCAAA	GAAAACGTGTC	TATGAAAATA	TTGCTTACGC	TATGGAAGTA	ATCGGGAAA	7260
ATCGCCGTA	TATCAAAAGA	CGAGTGATGG	AAGTTTGGGA	CTTGGTTGGA	TTGAAGCATA	7320
AGGTTCGTTC	TTTCCCAAT	GAACCTCTAG	GTGGGGAGCA	ACAGCGGATT	GGGATTGCGC	7380
GTGCAATTGT	AAATAATCCC	AAAGTATTGA	TAGCTGATGA	GCCAACAGGA	AATCTGGATC	7440
CGGATAATTG	ATGGGAAATT	ATGAATCTCT	TGGAACGGAT	TAACyTACAA	GGAACAACTA	7500
TTTTGATGGC	GACTCATATA	AGCCAGATTG	AAATACCTT	GGGCCACCGT	GTCATTGCCA	7560
TTGAAAATGG	CCGTGTCGTT	CGTGACGAAT	CAAAAGGAGA	GTATGGATAC	GATGATTAGT	7620
AGATTTTTTC	GCCATTATT	TGAAGCCTTA	AAAAGTTGA	AACGAAATGG	TTGGATGACA	7680
GTAGCTGCTG	TCACTTCAGT	CATGATTACT	TTGACCTTG	TGGCAATATT	TGCATCTGTT	7740
ATTTTCAATA	CAGCGAAACT	AGCTACAGAT	ATTGAAAATA	ATGTCCTG	AGTAGTTAT	7800
ATCCGAAAGG	ATGTGGAAGA	TAATAGTCAG	ACAATTGAAA	AAGAAGGTCA	AACTGTTACA	7860
AATAATGACT	ACCACAAGGT	ATATGATTCT	TTGAAGAAC	TGTCTACGGT	AAAAAGTGT	7920
ACCTTTCAA	GTAAAAGAAGA	ACAATATGAA	AAATTAACCG	AGATAATGGG	AGATAACTGG	7980
AAAATCTTTG	AAGGAGATGC	CAATCCTCTC	TATGATGCCT	ATATTGAGA	GGCAAACACT	8040
CCAAATGATG	AAAAAAACTAT	AGCCGAAGAT	GCTAAAAAAA	TTGAAGGTGT	CTCTGAGGTT	8100
CAAGATGGCG	GTGCCAATAC	AGAAAGACTC	TTCAAGTTAG	CTTCATTTAT	CCGTGTTGG	8160
GGACTAGGGA	TTGCTGCTTT	GTAAATTTT	ATCGCAGTTT	TCTTGATTTC	AAATACCATT	8220
CGTATTACCA	TTATTTCCCG	CAGTCGCGAA	ATTCAAATCA	TGCGCTTGGT	CGGAGCTAAA	8280
AACAGTTATA	TCCGTGGACC	GTTCTTGT	GAAGGAGCCT	TTATCGGTTT	ATTGGGAGCT	8340

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ATCGCACCAT CTGTTTGTT CTTTATTGTT TATCAAATTG TTTACCAATC TGTCAACAAA	8400
TCGTTGGTAG GGCAGAAATCT ATCCATGATT AGTCCAGATT TATTTAGTCC GTTGATGATT	8460
GCCTACTAT TTGTGATTGG GGTTTCATT GGTCATTGG GATCAGGAAT ATCCATGCC	8520
CGATTCTTGA AGATTTAGGT AAAATAGCTG CTTTATGAG GAGATTGTA AATCTCTTT	8580
TTTGCTACAA GAGTTTTGA AAAGAGATGC GCAGAAGAAA AGAGCTTCCA AAGAAGTCCC	8640
CCAGAGAAGA CTTC	8654

(2) INFORMATION FOR SEQ ID NO: 99:

- (i) SEQUENCE CHARACTERISTICS:
- (A) LENGTH: 19718 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 99:

TGTCGGGTCA AAATCATTAC TATGGCTATG TATAGCCCTT ACTATGACTT GGCTAACAC	60
GTTCGCTTTC AAATTTCTAG GCTCAGGCTG AAACAGTCTC CCAGGCTGTT CACTCCGAA	120
TGCTAAATC GTTCTTGATC GCTTCACAT TGTACAACAT CTTAGCCGTG CTATGAGTCG	180
TGTGCATGTC CAAATCATGA ATCAGTTCA TCGAAAATCC CATGAATACA AGGCTATCAA	240
GCGCTACTGG AAACTCATTC AACAGGATAG CCGTAAACTG AGTGATAAGC GATTTTATCG	300
CCCTACTTTT CGCATGCCT TAACAAATAA AGAAAATTCTT GACAAGATTG TAAGCTATTC	360
AGAAGACTTG AAACACCCT ATCAGATCTA TCAACTCTTA CTTTTCACT TTCAGAACAA	420
AGACCCGTGAG AAATTTTCG GACTCATTGA GGACAATCTG AAGCAGGTTTC ATCCTTTTT	480
TCAGACTGTC TTTAAAACCT TTCTCAAAGA TAAAGAAAAG ATTATCAACG CCCTCAACT	540
ACACTATTCT AATGCCAAC TGGAAGCGAC CAATAATCTC ATCAAACCTA TCAAGCGAA	600
TGCCCTTGTT TTTCGAAACT TTGAAAACCTT CAAAAAACGG ATTTTTATCG CTTTGAACAT	660
CAAAAAAGAA AGGACGAAAT TTGTCTTTTC TCGAGCTTAG CTGACTTCAA CCCACTACAG	720
TTGACAAAGA GCCTAATTTC CATAAAAATT GACATGGAAA TTATAAAACC ATTACTAGTT	780
TAGTCCTTT TGATAACGTG CCAATTGGC TTGGTTGCC CAAACATAGT GACCTGGACG	840
GATTCTTACG ATAGATGGCT TATCAGTCTC ATAGTCGTGT TGACTTGGAT CGTAAACCTT	900
CAAGACCTTC TTACGTTCCA AGATTGGATC TGGGATTGGT ACCGCTGAAA GCAAGGCTTG	960
AGTATATGGG TGAATTGGAT TGTAAACAA TTCTTCTGTT TCTGCAACCT CTACAATAAC	1020

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ACCCTTGTAA	ATAACTGCGA	TACGATCTGA	AATAAAGCGA	ACAACCGACA	AGTCATGGC	1080
GATGAAGAGA	TAGGTCAGGC	CGAGCTCTT	TTGGAATT	TTGAGCAAGT	TCAAGACTTG	1140
GGCACGTACA	GAAACGTCCA	AGGCTGAAAT	TGGCTCATCT	GCAATAACAA	AGTCTGGTTG	1200
CATGACCAAG	GCACGGGCAA	TACCGATACG	TTGACGTTGA	CCGCCTGAGA	ATTCATGAGG	1260
GTAACGAGTC	AACTGCTCAG	CAAGAAGACC	TACTTCACGG	ATAATATTTT	GAACCTTCTC	1320
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ATCAACAGTC	GCACGTTCAT	TCAAACATTGC	GGCAGGGTCT	TGGAAAATCA	TCTGGATTG	1440
ACGAATCAAT	TCCGCAGCCTT	GTTCACGCGA	TTTCTTACCA	TTAATCTTTT	GACCATCAA	1500
AATGTTATCT	CCATTACTTG	TATCATTAG	ACCGATGATA	GCACGACCAA	TAGTTGTTT	1560
CCCACCTACCG	GACTCACCTA	CAAGCGAGAA	AGTTTCTCC	TTGTTGATAA	AGAAGTTAGC	1620
ATTTTTAACCC	GGCACAAACT	TCTTACTTCC	TTCACCGAAG	GAATTTCTA	AATCTTTGAT	1680
TTCTACTAAT	TTTCAGACA	TTTCCTTCCT	CCTAGTCAGC	CAGATGGCA	AATCCCATT	1740
TTTCACGGAT	CTTATCATGG	AGATTTGCAA	TCACAGCTGG	TTTTTCTACT	TTCGGAGCAT	1800
CCTCATGAAG	AAGCCAAGTT	TTAGCCCAAT	GTGTCCTGTA	TACTGAGAAT	TGAGGAGCTT	1860
TTTGTTCGAA	GTCAATCTGC	ATTGCGTAGT	CAGAACGCAA	GGCAAAAGCA	TCCCCTTCA	1920
GGTCAGTATA	AAAGTACGGA	GGTGTTCCTG	GGATTGAGTA	AAGATCCCCT	TTATCATCAG	1980
CAAGCTGAGG	CAAGCTAGAC	AAGAGACTCC	ATGTATATGG	ATGGCGAGGG	TCATAGAAGA	2040
CTTCCTCAAC	CGTTCCATAC	TCAACGATTT	CTCCTGCATA	CATAACCGCT	ACCTTATCCG	2100
CAAACTTGC	CACCAACCCA	AGGTCTGGG	TAATAAAGAT	TGTTGTGAAA	TGATACTCGT	2160
TTTGTAAGA	TTTTAGCAAA	TCAATAATCT	GAGCTGAAAT	AGTTACATCC	AAGGCAGTTG	2220
TTGGCTCATC	ACAGATCAAG	ACATCAGGTC	GGCAGGCAAG	GGCAATAGCA	ATAACGATAC	2280
GTTGACGCAT	TCCTCCAGAA	TATTGGAATG	GGTATTCTATT	AAAACGTCTA	TCTGCGCTCG	2340
GAATGCCAAC	CTTATTCATG	TAGTCAATGG	CCAATTCTTT	CGCTTCTTTA	GCTGTTTTC	2400
CTTGGTGTGTT	TACAATAACT	TCTGTAATCT	GAETACCAAT	TGTTTTAATG	GGGTCCAAAC	2460
TAGTCATTGG	GTCTCTGGAAAG	ATAGTCGCAA	TCTTAGCACC	ACGAATTG	TCCCAATCCT	2520
TGTGAGAAGA	TAAAGCTGTC	AAAGTCTGAC	CACGGTAGTC	AATACTACCT	TGGGCAATAC	2580
GACCATTTC	TTCGAGCATA	CCTGTGAAGG	TCTTTGTCAA	AACAGATT	CCTGATCCTG	2640
ACTCACCTAC	CAAGGCTAAT	ACTTCTCCTT	CGACTAGTTC	AAGGGAAACG	CCGCGAATGG	2700
CTGTCAATAC	TTTGTACGAA	ACGTCAAATT	CCACGACAAT	ATCGCGAGCA	GTCAAAATT	2760
CATTTTTTTC	TTTGTCTATT	TCTACTCCTA	TCTATGTGTA	CGTGGATCAC	TAGCATCCGC	2820

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TAAGTTTGA CCAACTACGA AAAGGGACAA GGATACCAAG ACAAGGGTTG TCAATGGAAT	2880
CCAGAACAAAG TAAGCATTGG TTGTACGTT TTGTGAATAA TCCGAATCA AACGACCCAA	2940
ACTTGGCACT GAAATCGTA ATCCAAGACC GAAGAAAAGAC AAGAAGGCTT CGTATGAGAT	3000
AAAGCTTGGGA AGCATTGAG TCATGGTGT CACAATAACA GATACCAATT GAGGCATGAT	3060
ATTTTGGCA ACAATCTTCA AGGTTGGTGT TCCCAAAGTA CGTGACGCCA AGTTGTATTG	3120
CAAGTCACGA TAGCGCAAGA TTTGCACACG GATCATGAAG GCAATACCAA TCCATGTTGT	3180
TACGCTCATG GCAAAAATCA GATTCCAGAA TCCAGCTCCG ATTGAGTAAG TCAAGACAAT	3240
AACAAATCAAA AGAGGTGGGA TGTTTGAGAT GACGTTGTAA ACTTCATCA TGACACGGTC	3300
AACTGATTTT GAAATACCCC AAATACCACC GACAAAAACA CCGATAACCA AGTTAACAC	3360
TGTCGCAATC ACAGAAAATGA GGATGGAGTT ACGAGCTCCG AACCGACAC CGTCAAAGAG	3420
CGATTTACCG TTACTGTCAG TACCGAACCA ATGCTCCGCA TTTGGCTTGA TATAACGAAC	3480
ACTAAAGTCG TTTACCTTGC TGACATCATT GAAATCAAAC TTAGAAAACA TTGGGTAGAT	3540
GAAACTTATC AAAATGATGG CTACCAAGAT TCCCAACATG ACTACAGTTG ATTTTTCTT	3600
CATAAATTGT TAAACACTG ATTTCCAGTA AGAATATGCT GGCACATCAA TAGTTTCAGA	3660
GGCAAAATCG TCACGTTTA CAAACTGAAA TTTTCTTTA TCGATTGTAG ACATTATTTG	3720
CCTCCCTTCT CAGTCAATTAA AATACGTGGG TCAATAATAG TCATCCAAAT ATCTCCAAA	3780
AGACGTGAGA AGATAGAAAT ACATGTAAG ATGAAGACAA GACCAACGAC CATAGAGTTA	3840
TTAGATGCTT TTACAGAGTC AATCAACATT TTACCCATAC CTGGGAAGGC GAAGACTGTT	3900
TCAGTAAGGG TTGCACCAACC GATAACCCCCA ATAATGGCAG CAGGAATTCC TGAAACCAAGC	3960
CGAACCATGG CATTTTAAA GATGTGTTTG TTTGAAATTCTT CTTTTCTAGA CAAACCTTT	4020
GCACGAGCGA AACGAACAAA GTCTTGAGAT TGCAAGTCAA TCATGTAACG ACGAATCCAA	4080
ATGGCTGTAC CAGGAGCACCA CAACAAACCA AGGATGACTG CTGGTAAAC GTAAGAACGC	4140
CAATCTCCAG CTCCCAAGAT AGGGATGAA TCTGGAAAGGG CAATAGATGA TCCAATCAAT	4200
CGAACGATGT AAACCAAGGC AATCGTTGGAGA AGAGCAAGCA AGAAGGTCAA AGCCCTGTT	4260
GAGAGGCTAT CAATCCAAGT GTTCTTGAAA CGAGCCATGG CTGAACCAAG TGGCACGGCA	4320
AGAGCATAGG CAAGAACCAA ACCAATCAAAC CCAGTAATAG CAGAGCTGAC AATCATAGAT	4380
GGATATTGGT AATTACTTTC AGTCGCTGTA TAAGGATCAT CTTTCCATA GCTAGCTACT	4440
TCACGAGAGT CAGCCTGACT AGGTGACTTG TAGGTTCTTG ACTAAATATT TACAGAACAC	4500
GTTTTCTTAC CTGTTGGAA CTGAACCTGG GCAGTTTGTTGG TTTGTCCCTTG ACCTTGAGTA	4560

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ATAACCTGAA	GAACCTGGTGT	ATTAGCATAG	GTTGGGTAAG	AGTCACCTAA	ATTCAAGTTC	4620
ACAAAGTTTT	GATGAACAAA	TGGGAACTGA	CTGTTAAAGT	ACAAGAGATA	TTTATGTTA	4680
GTTCCCTGAAAC	CGACCAATGA	CCATCCGATA	GCTGGATCAT	TTTCAAACG	AAGGTAGCGT	4740
TTCAAGTCTG	GATTTTCAGG	GTCTTGATT	TTATTTGTAT	GGTCAATGTC	AATCAAGTTA	4800
GCATAGAAGT	GAAAAACACG	TTCAAAAATT	GGAATTTCAC	GAGTAGCATA	GAATTGACCA	4860
CTTCAGTAA	ATTCTCCCAA	AGTCCAACCA	TGACCTAATT	GATTGATGTA	CTTTTCATAA	4920
ATAGCTTAT	TGGTCGCATT	TGCTTCTACT	GTTACAGAAG	AATCCATGCT	ACTTGCCCTT	4980
TCTTGCAACT	CTTCTAGTATC	GTAATACTCA	ATGTAGCCCA	TACGCTCAA	CACAGTATTT	5040
TCATAGTTAT	CACGTTTATC	AGCCGTTGTC	GCAATTTTAT	TATAGTTAGG	ATCCTGCTTG	5100
AAAATCAATT	TTCGAGGAAC	CAAGGTATAG	ATAATCGTGT	AGGTCAAAGT	CGTTACTAAG	5160
AAAATCGAAA	CCAATGACCG	CAAACACGC	ATAAAAATAT	ATTTTTTCAT	ATTATTCCT	5220
TTAAAAATCC	CAAAGAAC	TTCTCCTCAT	GGAGAGAAAG	TTCTATTAGA	AATTATTTAC	5280
TTCACATGAC	TTGCCAATTC	TTTTTGAGCT	TTCTCATTTG	ATTCAGCTTT	TTCTTCAAC	5340
CATTTTCAC	GAGCTTTTC	ATACTCTTCC	TTAGTCACCA	CTTTATCTTG	TGATTTCAA	5400
TATTTGAAGT	AAACATCTGA	CCCCTTAGAG	CCTGTTGCG	CAGAAGCTCC	AGTAAATGGA	5460
ACAATTCGTG	AAAGCACTGG	TGCTGCACCA	GAAGAAGCCA	TAGCAGGAAT	AAAGAGTGAA	5520
CTATCTGTCA	ACCATGCTTG	AGCCGCTGCA	TATTTTCAT	AACGGACATT	CAAGTCGCTT	5580
GTCTCTCTGG	CAGCTTCATC	AACTAATTAA	TCGTATTCTT	TCAAACCAAC	TTGAACTACT	5640
GAAGGGCTAT	TTGGATTATC	AAATCCTAAA	TATGTTTTG	TAGTTTCACT	GCTAGTTGTT	5700
TTTAAAATAT	CCAGGTAAGT	AGATGGGTCT	TGATAGTCTG	GCCCCCATGA	AACTCCTCCT	5760
GATACATCCC	AATCCTCAGA	TGAAGCATTG	GCAGCATAGT	AAGTAATATT	AAGGAATTCA	5820
TCACCTGTCA	TTTGTGAAAT	ATCAACAACG	ACATTTCAA	CACCAAGAAC	TGTTTCTACA	5880
GATTGTTAA	AGGACTGAAT	ACGAGATATG	TAGTTTTTG	ATGCTGGTC	TACTGGAACG	5940
TCCAGATGAA	TAGGAAACTG	AACGCCGTCT	GCTTCTAAAG	CTTCTCTAGC	TTTCGCAAAC	6000
TCTGCCTTGG	CCTTGTCAAGC	ATTGAATAAA	CCATCCTGCC	CATCAGCTAA	ATTCACACCT	6060
TTCCACTCAT	CACCATAAAGC	AGGAAGTTGA	GCAGCGACTA	AATCACAAA	GGCTCTCTCA	6120
CCAGCTGAAA	CAAAGTCTGG	TTTACAAAT	AAATTACGAA	CTGCTAAAGC	TGCTCCATCT	6180
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AAATCTTGT	TAAGCAATGC	CTTCTTAGTA	GCTACTTTCT	CTGAATCTGT	AGTTTTAGAA	6300
GTATAGTTGT	AACTTTGGCG	ATCAATATTC	ACACCCAGAC	CAGCAATCCC	AGAGCCTGAT	6360

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TGTGTGTAAT AGATATTGTC CTTGTATTCT TCTGCAACCT TAGAATAGTT GGAGCTGGTA	6420
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TCTGATCCAT CATAGTAAGC TAGATTGATA GTATCTAGGT GGACATTTTC TTTATCCCAA	6540
TATTGCTCAT TTTTTACAAA CTCTACAGAA GATTTGCAG TCAACCCCTT CAACAAGAAC	6600
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GCATGTTCAT GATAAGGAGC ATGTATCCGA GGATAAGAGT TCATATAGGT CAATGGCCCT	8280
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TTGGCATAAC	CGAATTGGTT	GTTTTTAAAG	GCATAGTTAT	AAACCAAGAG	CCCAAGTGAG	12060
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CCACCTTTTA	GGGCTAGGAT	AAAGACCATA	GAGACACTTG	GTAGCAAGTA	AGGCAATTCA	12180
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GACATCCCCA	CGTGCAGAAGG	CTCCGATAAC	ATCGGTATAG	CCAGCACCTT	CCCAGTTCTT	13320
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TATCCAGTTG AGCTAAGGGT GCTCCATATT ATGCCGAGGA CGCGAATCGA ACCGGTACGA		19260
TCGTTACCAA TCGCAGGATT TTAAGTCCCTG TGCGTCTGCC AGTTCCGCCA CCCCCGGCTC		19320
TCTAAGCGAA CGACGGGATT CGAACCCCGCG ACCCCCCACCT TGGCAAGGTG GTGTTCTACC		19380
ACTGAACTAC GTTCGCACTG TTTTCTCTA TCTAAAAATG CCGGCTACAT GACTTGAACA		19440
CGCGACCCCTC TGATTACAAA TCAGATGCTC TACCAACTGA GCTAAGCCGG CTCAATTGTT		19500
ATATCTTAAT GCGGGTTAAC GGACTTGAAC CCCCCACGCCG TTAAGCGCCA GATCCTAAAT		19560
CTGGTGCGTC TGCCAATTCC GCCAAACCCCG CATATATGAC CCGTACTGGG CTCGAACCAG		19620
TGACCCATTG ATTAAAAGTC AATTGCTCTA CCAACTGAGC TAACCGAGTCT AAAATAACTT		19680
GCGTACCTT AAACGGTCCG ACGGAAATCGA CCCCCGGTAC		19718

(2) INFORMATION FOR SEQ ID NO: 100:

- (i) SEQUENCE CHARACTERISTICS:

 - (A) LENGTH: 4117 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 100:

CCGTGGAAAA	GTCTGGATAG	TGAATGGTCT	TCACACAATG	ACCTGAAAGA	AGCCTGAGAA	60
TAATTATGGA	GAGTAGCATT	CTGAGAGGTG	TTAGCAGAAC	CATATGACAG	AGCTGTTTGA	120
AGAGGGAAATA	TTGAGGGAGAA	AAATCCTGAG	CCTACCAGTT	GGACTTGGAA	AGAGCTGACT	180
GTTAGATCAT	GGTTTATTAT	CCACAACTG	TGGATAACTT	TGTGAATAAG	AGAAGTTGCT	240
AAAGAAGGAG	ATATATAACG	ATGAAGAAAA	TCAAACCGCA	TGGACCGTTA	CCAAGTCAGA	300
CTCACTAGC	TTATCTGGGA	GATGAACTAG	CAGCTTTAT	CCACTTCGGT	CCTAAATACCT	360
TTTATGACCA	AGAATGGGGG	ACTGGCACAGG	AGGATCCTGA	GCGCTTTAAC	CCGAGTCAGT	420

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TGGATGCGCG	TGAGTGGGTT	CGTGTGCTCA	AGGAAACGGG	CTTCAAAAAG	TTGATTTGG	480
TGGTCAAGCA	CCACGATGGC	TTTGCCTTT	ATCCGACAGC	TCACACAGAT	TATTCGGTTA	540
AGGTCACTCC	TTGGAGGAGA	GGAAAGGGCG	ACTTGCTCT	TGAAGTATCC	CAAGCTGCCA	600
CAGAGTTGA	TATGGATATG	GGGGTCTACC	TGTCACCGTG	GGATGCCCAT	AGTCCCCTCT	660
ATCATGTGGA	CCGAGAACGCG	GACTACAATG	CCTATTATCT	GGCTCAGTTG	AAGGAAATCT	720
TATCAAATCC	TAACTATGGG	AATGCTGGTA	AGTCGCTGA	GGTTTGGATG	GATGGTGCCA	780
GAGGAGAGGG	CGGCCAAAAG	GTAAATTATG	AATTTGAAAA	ATGGTTGAA	ACCATTCTG	840
ACCTGCAGGG	CGATTGCTTG	ATTTTTCAA	CAGAAGGCAC	CAGTATCCGC	TGGATTGGCA	900
ATGAACGAGG	GTATGCAGGT	GATCCACTGT	GGCAAAAGGT	GAATCCTGAT	AAACTAGGAA	960
CACAAGCAGA	GCTGAACATAT	CTTCAGCACG	GGGATCCCTC	GGGCACGATT	TTTTCAATCG	1020
GAGAGGCAGA	TGTTTCCATC	CGTCAGGCT	GGTTCTACCA	TGAGGATCAG	GATCCTAAGT	1080
CTCTCGAGGA	GTTGGTCGAA	ATCTACTTTC	ACTCAGTAGG	GCGAGGAAC	CCACTCTTGC	1140
TTAATATTC	GCCGAATCAA	GCTGGCTCT	TTGATGCAA	GGATATTGAA	CGACTTTATG	1200
AATTTGCGAC	CTATCGCAAT	GAGCTCTATA	AAGAAGATT	GGCTCTGGG	GCTGAGGTAT	1260
CTGGTCCAGC	TCTTCCGCA	GACTTTGCTT	GTGCCATT	GACAGACGGC	CTTGAGACCA	1320
GCTCTTGGC	AAGCGATGCA	GACTTCCCCA	TCCAGTTAGA	ACTCGACTTA	GGTTCTCCTA	1380
AAACTTTGA	TGTAATTGAG	TTAAGAGAAC	ATTGAGACT	AGGGCAACGA	ATCGCTGCTT	1440
TTCATGTGCA	AGTAGAGGTG	GATGGTGTCT	GGCAGGAGTT	TGGTCGGGT	CATACTGTTG	1500
GTTACAAACG	TCTCTTACGA	GGACGAGTTG	TTGAGGCACA	GAAGATACGT	GTAGTCATTA	1560
CAGAATCACA	GGCTTTGCCT	TTGTTGACCA	AGATTTCCCT	TTATAAAACT	CCTGGATTAT	1620
CAAAAAAAGA	AGTTGTTCA	GAACCTAGCAT	TTGCAGAAAA	AAGCCTAGCT	GTGGCAAAGG	1680
GAGAAAATGC	CTATTTACA	GTTAACCGCA	GAGAATGTAG	TGGCCTTTA	GAAGCTAAGA	1740
TTTCGATTCA	ACCGGGGACA	GGTGTCCATG	GTGTCGCCA	TCAGGATGAG	ATTCAAGTCC	1800
TTGCGTTCA	AACTGGTGAG	ACTGAAAAAA	GTCTGACGCT	ACCAACCTTG	TATTTCGCAG	1860
GAGATAAAAC	CTTGGATTTC	TATCTGAACC	TAACGGTGG	TGGTCAGCTT	GTGGATCAAC	1920
TTCAAGTCCA	AGTTTCATAA	AAGAAGAAC	TTTGCACGAT	GCAAAGGTT	TTTTGGTTAT	1980
TAGTGACTTG	GTAACCGACT	GAGGGTGAAA	GTAGTTGTT	CAGCTTTAA	GAGGTCTTGG	2040
TGTTGAATAG	TTGATACGAG	TGTTTGTCC	AGTCGGCATT	CTTTGACAAA	GTTAAAATGG	2100
TTGTGGTTTT	GTTTAGTATG	GATATCCAGC	CATTTATCTT	CTTACGCGAG	GTAGACTCGT	2160

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AGATGGTCAA AGAGAGGGAT TCCGAGGTCA TAGCTTGGTT TTCCCTGGACA GGTTGGATAA	2220
AATCCGAGAG CTGACCAGAT GTACCAAGCA GAGAGACTAC CATTGTCCTC ATCTCCAGGA	2280
TAGGCTTCCC AACTGGGTG AAAAGCTTTC TGACGGAGCG TCTTGATAAG AAGGGCAGTG	2340
TAGTCAGGGT AATCGCTGTA ACAGGAAGAGA TAAGGAATGT GGAAACTAGG CTGGTTGGAA	2400
ATGGCTATTT GTCCAAAAGG AGCAGTAGCC ATCTCGCTCA TTTCGTGAAT TTCGTAACCA	2460
TAGCCTGTTG TTTCAGGAGA GGGAGCATCT TGACAGGCTT TCAAAAGATA GTTGCTAAAG	2520
GTTTCTTTTC CACCCATCAG TTGGATTAAG CCAGGGATGT CGTGGAGAAC GCCTAAAGTA	2580
GCTTGAATGG CAGAGCATTG AGCGTAGTCT CGCCCCAAC TATAAGGAGA GAAGTCAGGG	2640
TGAAAGTTTC CTTGATTGTC TCGTGCTCGC ATGTAACCTG TCTCAGCGTC AAATAGCTGG	2700
CGGTAATTTT GTGAAGCAGC CTTGTAGGTT TCAGCGATTT CTATGTTCTC TAGTTTTTG	2760
GCACAGCTGG CGATACAAAA GTCACTATAG GCATAGTCTA GAGTATGGCT AACACTTCG	2820
TGGTGGTCGG TAGAGAGGTA ACCTAGTTCT TGGTATTGGG CTAGTCCGTG GCGGCCATTG	2880
ATGCCGAGAG GGTCGGCTTT GCTGGCTGTT TCGAGCATGG CTTGGAAGAG TTCTCCTTCT	2940
AGGTGGGGGG TCATGTCCTT GCAGCGCTA TCTCGGATAA TACCGCTAA AAGTGTACCT	3000
GGCATCATAC CCCGTTCATC TGGAGCCAGC CATTGGAA GGAAACCAGT ATCGCGGTAG	3060
CTATTGAGGA AACCTTCTAA AAAGCGTTGA TAGTGCTCCG GTATGATAAG GGCAAAGAGG	3120
GGGAAGGTGG TGCGGAAGGT ATCCAGAAA CCATTGTTGC TAAAGAGGAC ACCAGGCTTG	3180
ACAGTACCAAG TAGCCAGATC CATGTGGATG GCTTGCCCTG ATTCAATTAT CTCATAAAA	3240
GTCTGTGGGA AGAGGAAGAG TCTGTAGAGG CAGTGGTCAA AGAAGGTTCG GTCAGCCTCT	3300
CCTGTCCTCTA TAATGTCAAA ACAGATGGAGG AGATTTCCC AATCCACTTG GGCACATTGAT	3360
TTACAGCTAT CAAAATCTTC TTGAGGTAGA TTGATTAGAG CTTGAGAAGG AGAGATGAAA	3420
GAAGTGGCTA GTTCATCTC GGTGACTA CTTGCTAAAGT CAATTCGCCA GTCTCCAGCT	3480
TCTTGGCTGA TAGCAAGAAT ATCCGTGTTA ATTTCAGGG CAGTGAACAT CGTTAGCGAA	3540
TTTTTGTAG TTTCAAGTTT ACCTTCTGTG CGCAGGGCAA GAGTCCGCTT ATCTACTTGC	3600
TCTACTGTCA GTTCATCTGC TCGGTGAAGA TAGAGGGAGA GGGCTTGCC TTGCTTTGA	3660
TTCAAAACGAA TAGAAGCACC ATAGCAAGTC GGTGTGAGCT GGGTTCAAT CTGATAACGC	3720
AGAGAAAAGA GCTTCAAATA GTGAGGCTGG AAGCAAGCTT TATCTATATC ATAAGAAGAC	3780
TGGCGGTGAA AGAGGCTGTC TCCCCCAGT TGACTGGTGA CAGGTGTCAG AAGGAGCCAA	3840
GAGTAGTCCC CAATCCAAGG ACTGGGCTGG TGAGTTAACG GAATCCCTG AAAGATAGGC	3900
AGATGTGGAT CAAAAAACCA AGATCCATCC TGGTCACTGG TCTGGGGCAC AAAGTAATTC	3960

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ATCCCAAAAG GCACGCCTGT GTATGGCAGG GTATTTCCCC GAGAAAAGGC ATGCTTGTG	4020
GTAGTTCCAA AACGGGTATC GATGGTATCA AGTAGTGTT TCATAGTCTT TCCTTTAGCT	4080
GTTCCTCTAC ATTATATCAG TAATAGAGGG CCTTTAG	4117

(2) INFORMATION FOR SEQ ID NO: 101:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 2727 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 101:

CTGGTTCAAT TATTATTACAC TCTAAGTAGT CATATGTTCT TTATTTATGT GAGTTTTAC	60
CTTTAAAGG ATCTTGTAG ATGGGAGAAG GTTTTAAAG TGACAGATGA TAATACAAGA	120
AAAGTTCGTT TATTAGTAGC CTTTTTTAGC ATTGTATAG GCTACATCCT GAGTTCTTTC	180
TTTATTTAGCC TGTATCATTT GTGGCAAGAA GCGCTTAGAG GATTATTATG AAATCAAGAG	240
TAAAGGAAAC GAGTATGGAT AAAATTGTGG TTCAAGGTGG CGATAATCGT CTGGTAGGAA	300
GCGTGACGAT CGAGGGAGCA AAAATGCAG TCTTACCCCTT GTGGCAGCG ACTATTCTAG	360
CAAGTGAAGG AAAGACCGTC TTGCAGAATG TTCCGATTTT GTCGGATGTC TTTATTATGA	420
ATCAGGTAGT TGGTGGTTTG AATGCCAAGG TTGACTTTGA TGAGGAAGCT CATCTGTCA	480
AGGTGGATGC TACTGGCGAC ATCACTGAGG AAGCCCTTA CAAGTATGTC AGCAAGATGC	540
GCGCCTCCAT CGTTGTATTAA GGCCCAATCC TTGCCCCGTGT GGGTCATGCC AAGGTATCCA	600
TGCCAGGTGG TTGTACGATT GGTAGCCGTC CTATTGATCT TCATTTGAAA GGTCTGGAAG	660
CTATGGGGGT TAAGATTAGT CAGACAGCTG GTTACATCGA AGCCAAGGCA GAACGCTTGC	720
ATGGTGCTCA TATCTATATG GACTTTCCAA GTGTTGGTGC AACGCAGAAC TTGATGATGG	780
CAGCGACTCT GGCTGATGGG GTGACAGTGA TTGAGAATGC TCGCGCTGAG CCTGAGATTC	840
TTGACTTTAGC CATTCTCCCTT AATGAAATGG GAGCCAAGGT CAAAGGTCTT GGTACAGAGA	900
CTATAACCAT TACTGGTGTT GAGAAACCTTC ATGGTACGAC TCACAATGTA GTCCAAGACC	960
GTATCGAAGC AGGAACCTTT ATGGTAGCTG CTGCCATGAC TGGTGGTGAT GTCTTGATTC	1020
GAGACGCTGT CTGGGAGCAC AACCGTCCCT TGATTGCCAA GTTACTTGAA ATGGGTGTTG	1080
AAGTAATTGA AGAAGACGAA GGAATTCTGTG TTCTGTTCTCA ACTAGAAAAT CTAAAAGCTG	1140
TTCATGTGAA AACCTTGCCC CACCCAGGAT TTCCAACAGA TATGCAGGCT CAATTTACAG	1200

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CCTTGATGAC ACTTGCAAAA GCGAATCAA CCATGGTGG AACAGTTTC GAAAATCGTT	1260
TCCAACACCT AGAAGAGATG CGCCGCATGG GCTTGCATTC TGAGATTATC CGTGATACAG	1320
CTCGTATTGT TGGTGGACAG CCTTGCAGG GAGCAGAAAGT TCTTCAACT GACCTTCGTG	1380
CCAGTGCAGC CTTGATTTG ACAGGTTGG TAGCACAGGG AGAAACTGTG GTCGGTAAAT	1440
TGGTTCACTT GGATAGAGGT TACTACGGTT TCCATGAGAA GTTGGCGCAG CTAGGTGCTA	1500
AGATTCAAGCG GATTGAGGCA AGTGTGAAAG ATGAATAAGA AATCAAGCTA CGTAGTCAAG	1560
CGTTTACTTT TAGTCATCAT AGTACTGATT TTAGGTACTC TGGCTCTAGG AATCGGTTA	1620
ATGGTAGGTT ATGGAATCTT GGGCAAGGGT CAAGATCCAT GGGCTATCCT GTCTCCAGCA	1680
AAATGGCAGG AATTGATTCA TAAATTTACA GGAAATTAGG CTGGAGAAC AGCCTTTTC	1740
TAAAGATAAG GAGAAATATG AACAAAAAAA CAAGACAGAC ACTAATCGGA CTGCTAGTGT	1800
TATTGCTTT GTCTACAGGG AGCTATTATA TCAAGCAGAT GCCGTCGGCA CCTAATAGTC	1860
CCAAAACCAA TCTTAGTCAG AAAAACAAAG CGTCTGAAGC TCCTAGTCAA GCATTGGCAG	1920
AGAGTGTCTT AACAGACGCA GTCAAGAGTC AAATAAAGGG GAGTCTGGAG TGGAATGGCT	1980
CAGGTGCTTT TATCGTCAAT GGTAAATAAAA CAAATCTAGA TGCCAAGGTT TCAAGTAAGC	2040
CCTACGCTGA CAATAAAACA AAGACAGTGG GCAAGGAAAC TGTTCCAACC GTAGCTAATG	2100
CCCTCTTGTC TAAGGCCACT CGTCAGTACA AGAATCGTAA AGAAACTGGG AATGGTTCAA	2160
CTTCTTGAC TCCTCCAGGT TGGCATCAGG TCAAGAATCT AAAGGGCTCT TATACCCATG	2220
CAGTCGATAG AGGTCAATTG TTAGGCTATG CCTTAATCGG TGGTTGGAT GGTTTGATG	2280
CCTCAACAAG CAATCCTAAA AACATTGCTG TTCAGACAGC CTGGGCAAAT CAGGCACAAG	2340
CCGAGTATTC GACTGGTCAA AACTACTATG AAAGCAAGGT GCGTAAAGCC TTGGACCAAA	2400
ACAAGCGTGT CCGTTACCGT GTAACCCCTT ACTACGCTTC AAACGAGGAT TTAGTTCCCT	2460
CAGCTTCACA GATTGAAGGC AAGTCTTCGG ATGGAGAATT GGAATTCAAT GTTCTAGTTC	2520
CCAATGTTCA AAAGGGACTT CAACTGGATT ACCGAACCTGG AGAAGTAAC GTAACTCAAGT	2580
AAAAGATACG CCTACACTCC TATGTCACTT ATGGATGTAG GAGTTCTTT TACTAGTTA	2640
AGCAGGACTA AGACAGGTAC TAAGACAAAA TAGCAACTTC TAAAACTAAC TTCCAGTTT	2700
GGGAGAGAGA TGGAAGTTAC TTTGAGA	2727

(2) INFORMATION FOR SEQ ID NO: 102:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 5717 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 102:

TTTTTTGTAG ATTTAAGTGG GGTGCAATT CAAAAAAATA AAAAACATT TTTGAAAATT	60
ATGTTAGCAG GAATTGCTTC AAATTCGATT TTATCACTTA CAGGTTTACT TGTTTTATTG	120
TTCACATCGT ATAATATGCT TGGACTCTTA TTTTTTATCA TTAACTTAGG TATGATTTT	180
ATTAATTCAA TTCCCTTTTT TCAGTATGAT AGTGGTATTA TTTTAAGATA CTTGAATTCT	240
AACAATAATA ACTTGAATTT TCAATATATA GTTCAACTTT TAATAGCATT TGTTATTATT	300
TATTTTCCTT TGAGTCAACT ATTACAGTTT TTGACACCCA ATATTATTGT TCGTAGTATA	360
GGAGGGGTGG TTGTTTCTAT ACTGCTTTCT ATATTATATA TGATAGGAAG GACGAAATAT	420
GTTCTACGTA AATAGTTATG TTTTGCTTA TAAAAAAAGAA GGTATAATGT ATTTACGTGG	480
TCGGAGTATG CGGGAAATAG CTATAGAACCC TCAAATTCG CAAGAATTAA TCAACGATCT	540
ATTTAATAGT TCTAAGGAAC TATTAGAGAT AGAAGAAGTA TTAGGCAGTA AACTAACATT	600
TGAACTATAA ATGAACAAAT TTTAATTTCG GATGAGATAG ATATTGATAG TAGATATTCT	660
AGAACTAAAG GTTACTATTC GTTATTTTAT AATGAAGAGT ATAATAAAAT ACAGAATAAA	720
ACAGTATTAG TATTAGGAGC AGGAGTCTTA GGATGTTATA TATCTCTAAG TCTAAGTATG	780
TATGGAGTGA GGAAACTTAT TGCGCTGAT TACGATATAA TAGAACCATC AAATTTAAAT	840
AGGCAAATTC TTTATACAGA GTCGGATGTT GGTAAGGAGA AGATTAATGT TCTTTCTGAA	900
AAAATACACA AGTATAATTC AGATGTTCAAG GTAGTACCTA TTTCTATTAA AGTTTCTTCA	960
GTAGAAGAAT TAGAAAAAAAT TGTTGCGGAA TATGGGAGTA TAGATTTTAT CGTTAAAGCA	1020
ATTGATAACGC CCATTGATAT TATAAAAATT GTCAATCAAT TTGCTGTATC GCATAAGATA	1080
TCCTACATAT CAGGAGGGTT TAATGGATGCC TATCTTATTA TTGATAATAT ATATATCCCT	1140
ACCATCGGTT CTTGCTTTGG TTGTCGGAAT ATAACAAAG ATATAAATAA GTACACTTTA	1200
TCTGATAAGA CAAAGTGGCC GACTACACCA GAGATGCCTG CTATTTGGG AGGGATAATG	1260
ACTAATTTAA TAATTTAAAT ATTTCTGGGA TGTTATAATG AAATCCTAAT AGATAACGCT	1320
TACGTTTATA ATATGAGAAA TCATGCTCTA AGTCAAGAAA AATATGTTCT GGAAACGGA	1380
GAATGTCCAA TTGTAAGAA AATAATAAAG TGAAAGATAA CAATATTAGA GCGAAAACAT	1440
TTATTCGTTCA AGTTTGTGTT TGCTTATTAT CAGGAGGAGT AGCTTTTTA TCTGCTATTG	1500
GGCAGTTCAC TGTTATAGAA ACACAATTAA TAGTATTGTT CTTGGGTATT ATTTTTGCTA	1560
TATATTATGC TTACTACAAT AAAATATTC AAACATCATT GGAAATATA GTATGGCTTT	1620

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TTTCATCGTT	TGAGATTATA	TTTTGCTTG	TTAATTTTAG	AACATTTATT	CAGTTACCA	1680
TGGATATTTT	TATTGGTATG	ATAATATTTT	TAATGCTGTG	GATATTTATT	ATGTTAGGTA	1740
TAGTGTGTCT	TAGTATTAT	ATAACTTTAT	TATTTAGCAA	GGAGGCTTAG	TATGTTAA	1800
AAAATAGGTA	TAATGAGCAT	TTGCATATAT	ATAATTATTT	TATACTGCTT	GAGAATGTAT	1860
CGTATTATCA	ATAATATTGA	AACAATCTG	CTAACGGTTA	TATGCTTAAT	GTTATTGTT	1920
TTTTAAGAC	GTAAAGATAAG	AAAAGATAAG	TTAAAGATG	TTAAAGTAAA	ATGAGAATA	1980
TAAAGGAGGT	GCAATGAGTA	TGATTGAAGT	TAGCCATTAA	TCAAAAGTT	TTGGTGATAA	2040
AATAGCTTTA	AATAATATAA	GCTTCACTGT	TTAAAGAAGGT	TAGATTTTG	GATTTTAA	2100
ACCATCTGGT	TCTGGAAAGA	CCACAACGAT	TAATATTCTG	ACTGGCAGT	TCCTTGCCGA	2160
TAAAGGACAA	TCTATTATTT	TGGGACAAAA	ATCTCAAAAT	TTAACAAAGCG	GTGAATTAA	2220
GAGAATTGGA	TTGGGTTAGCG	ATACAAGTGG	ATTTTATGAG	AAAATGTCTC	TGTATAACAA	2280
TCTTCTTTT	TATAGTAAAT	TTTATAATAT	TAGTAAATCA	CGTGTGATA	ATTTGTTAA	2340
GCGAGTAGGA	TTATATGATA	GTCGCAAGAT	GGTAGCAGGA	AAATTATCCA	CTGGAATGAG	2400
GCAACGAATG	CTTTTAGCAC	GAGCTCTTAT	CAACAACCCC	GCTGTACTCT	TTCTGGATGA	2460
ACCGACCTCA	GGTCTAGATC	CCACAACCTTC	TCGAACAATT	CATGAGTTAA	TTTTAGAATT	2520
GAAAACAGCA	GGGACAACGA	TTTTCTAAC	GACTCATGAT	ATGAATGAAG	CAACTCTTT	2580
ATGTGATTAT	GTTGCCATTAT	TAATAAAAGG	GAAATTAGTT	GAGCAAGGAG	CTCCTTCTGA	2640
ACTCATTCAA	AGATATAATA	AAGATAAAA	GATTAAGGTT	ACAGATTATA	ATGGAATCA	2700
GATAACTTTT	GATTTTACAT	CACTAGAACCA	GGTATCTCAG	ACTGATCTGG	AAAATATTTT	2760
TTCAATTCA	TCATGTGAGC	CTACTTTAGA	AGATATTTTT	ATCACATTAA	CAGGAGGAAA	2820
GCTAAATGCT	TAACGGTTT	CTGGCTTTGG	TATGGTTGCG	TTGTCAAATC	ATCCTTCCA	2880
ATAAGAGTAT	TTTATTGCAA	GTGTTAGTGC	CTTTTGCTTT	CACATATTTT	TATAAATATC	2940
TTATGGAAAC	ACAGGGGAAG	GTCACACGATC	AACAGGCATT	AGTTCTTTG	ATGATGTGTT	3000
TACCTTTTC	TTTTCTTTG	GCTGTTGGAA	GTCCTATAAC	TATTATCTG	TCTGAAGAAA	3060
AAGAAAAGTA	CAATTACAA	ACTCTCTGT	TGAGTGGTGT	AAAAGGCTCC	GAATACATT	3120
TATCAACTAT	GTTCCTTCCT	TTTTGCTAA	CTTTTGTGAT	TATGGGAACT	ACTCCTCTTA	3180
TTTTAGGAGT	TACAATTGTA	CATACTTTA	ATTATATTAC	AATCGTTCTT	CTAACCTCTT	3240
TATCCATCAT	TTTATTCTAT	TTATTGATAG	GTPTAACCGC	GAAGAGCCAA	GTAGTAGCTC	3300
AGGTTATCAG	TCTTCCTGCT	ATGATTTAG	TTGCTTTCTT	ACCGATGCTA	TCTGGTTGG	3360
ATAAGACAGT	TGCGAAGATA	ACAGATTATA	GTTTATGGG	ACTATTTACT	AAGTTTTCA	3420

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ACTCGTACCC TAATTGAAAA TGGTTTGTC GAAGCTATCA AAGAAGCCTT TCCTGAAGTA	5220
GAAGTGATTG CAGGAACCTGC AACAGCAGGG ATTCCACACG GAGCCATTAT TGCTGATAAG	5280
ATGGACTTGC CTTTGCCTA CATCCGTAGT AAACCAAAAG ACCACGGAGC TGGTAATCAA	5340
ATCGAAGGTC GCGTAGCTCA AGGTAAAAA ATGGTAGTGG TTGAAGACCT TATTTCAACG	5400
GGTGGTTCAAG TTCTTGAAGC TGTAGCAGCA GCCAAGCGAG AAGGAGCAGA TGTACTTGGA	5460
GTTGTAGCGA TTTTCAGCTA CCAATTGCCA AAAGCAGATA AGAATTTCG AGATGCTGGT	5520
GTTAAACTTG TGACGCTTTC AAACATATAGC GAGCTTATCC ATCTAGCCCA AGAAGAAGGT	5580
TACATCACCG CAGAGGGCCT TGATCTTCTA AAACGCTTTA AAGAAGACCA AGAAAATTGG	5640
CAAGAAGGTT AGGTCAAGTA GATAAAGAGA GACGAGGCTA CCGAGTCTCT TTTACCATT	5700
TATTTAAAAAT ATGACAG	5717

(2) INFORMATION FOR SEQ ID NO: 103:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 5558 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 103:

CCTGGACTTT CTAAAATGAA ATCTTGCAC CTGGATCAAG CCCTTCATGA GCATTTTTCA	60
GAAGAAGAAT TAGCTGGTCA CTTTCATGTC CTTCTATGGA CTTTTTTTAC AATGGCATTG	120
CTATCACACC CAATACCTAT CTAAGGCCCT GGTCGTAAA CTTTATTGCA GCTCTTCCTC	180
TAAATTTCCT AATTGTTGAA CCAATTGCCC GTTTTATACT AAGTTCTTTT CAGAAACCAT	240
TTACTGGGGA AGAAGTTGAA GATTTCAAG ATGATGATGA AATCCAACCT ATTATCTAAG	300
CCAGTCTGTAA AACTACTAA TATTTGAAAT CCACTTCCTT TTAGGGTGCA ATGGTTATAA	360
ATGAATTTTT GAGAGGATCA GAATGAAAAA ACTAGCAACC CTTCTTTTAC TGTCTACTGT	420
AGCCCTAGCT GGGTAGCA GCGTCCAACG CAGTCTGCGT GGTGATGATT ATGTTGATTC	480
CAGTCTGCT GCTGAAGAAA GTTCCAAAGT AGCTGCCAA TCTGCCAAGG AGTTAACGAA	540
TGCTTTAACAA AACGAAAAGC CCAATTCCCC ACAACTATCT AAGGAAGTTG CTGAAGATGA	600
AGCCGAAGTG ATTTTCCACA CAAGCCAAGG TGATATTGCG ATTAAACTCT TCCCTAAACT	660
CGCTCCTCTA CGGGTTGAAA ATTTCCTCAC TCACGCCAA GAAGGCTACT ATAACGGTAT	720
TACCTTCCAC CGTGTACATCG ATGGCTTTAT GGTCCAAACT GGAGATCCAA AAGGGGACGG	780
TACAGGTGGT CAGTCCATCT GGCATGACAA GGATAAGACT AAAGACAAAG GAACTGGTT	840

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CAAGAACGAG ATTACTCCTT ATTTGTATAA CATCCGTGGT GCTCTTGCTA TGGCTAATAC 900
TGGTCAACCA AACACCAATG GCAGGCCAGTT CTTCATCAAC CAAAACCTTA CAGATACCTC 960
TTCTAAACTC CCTACAAAGCA AGTATCCACA GAAAATTATT GAAGCCTACA AAGAAGGTGG 1020
AAACCCTAGT CTAGATGGCA AACACCCAGT CTTTGGTCAA GTGATTGACG GTATGGATGT 1080
TGTGGATAAG ATTGCTAAGG CGAAAAAGA TGAAAAGAC AAGCCAACTA CTGCTATCAC 1140
AATCGACAGC ATCGAAGTGG TGAAAGACTA CGATTTAAA TCTTAAAAAC CAAAAAAATA 1200
CAGTATCCAC ATTCCGTACT GTATTTCTTT TACTCTCATT CTTAAGTTAA ATTATTTAAA 1260
TCCCCATATTT GGTCTATCCA GCCTTCATAA AAGTCTGGCT CGTGGCAGAC CATAAGGATA 1320
GATCCCCAT ATTCTTTGAG AGCCGCCTTG AGCTCATCCT TTGCATCCAC ATCCAAATGG 1380
TTGGTCGGCT CGTCCAGCAC TAAAACGTTG TTTTCACGAT TCATCAAGAG ACAGAAACGA 1440
ACCTTGGCTT GCTCTCCCCC TGATAATACT TGAATCTGGC TTTCAATATG TTTGGTTGTC 1500
AAACCACAAAC GGGCAAGGGC TGACCGACT TCTGCTTGAT TAAGGGCAGG AAAGGCATTC 1560
CAGACAGCTT CAAGAGGAGT TTGGCGATTA CGGCCTTCTA CTTCCGTGTC AAAATAACCA 1620
AGTTCTAAAT AATCTCCACG CTCCACTTCC CCAGCGATTG GCGAGATAAT GCCCAAGAGA 1680
CTCTTCAGA GAGTTGTTT TCCAATACCA TTAGCACCAA TAATCGCAAC CTTTTGATTG 1740
CGTTTCGAAGG TAAGATTTAA AGGCTTAGTA AGAGGACGGT CGTAACCAAT TTGCAAGTTC 1800
TTGGCTTGGGA AGATAAAAGCG CCCTGGTGTGTA CGAGCTGGTT TGAAATCAAA GGATGGTTTT 1860
GGTTTCTCAC TTTGGAGTTC GATAATATCC ATCTTATCCA ATTTCTTTG ACGAGACATA 1920
GCCATATTAC GAGTTGCAAC ACGGGCTTTA TTACGAGCCA CAAAGTCCCT GAGGTCTGCA 1980
ATCTCTTCT GCTGGCGTTC GTAGGCTGCC TCTAGCTGAG ATTTCTTCAT AGCATAAACT 2040
TCTTGGAACT GGTAGTAGTC ACCAGAGTAA CGCGTCAGCT GTTGATTTTC CACATGATAG 2100
ACAATATTAA TAACGTCATT GAGGAATGGA ATATCGTGCN AAATGAGAAC AAAGGCATTC 2160
TCATAGTTT GGAGATAGCG CTTGAGCCAA TCAATATGCT CAGCATCCAA GTAGTTGGTC 2220
GGCTCGTCCA ACAGCAAGAT ATCAGGCTTT TCAAGGAGAA GTTTTGCCAA AAGCACCTTG 2280
GTTCTTGCC CACCTGACAA AGAAGTTACA TCCGTATCCA TGCCAAAGTC CATAACACCA 2340
AGAGCACGCG CTACTTCGTC AATCTTAGCA TCCAAGGTAT AGAAATCACG ACTCTCCAGA 2400
CGGTCTTGAA GTTCTCCTAC TTCTTCCATG AGAGCATCAA CATCCGCGCC GTCTTCAGCC 2460
ATTTTCATAT AGAGGTCACT GATACGAGCT TCAGCTTTGA AAAGCTCATC AAAAGCCGTA 2520
CGGAGAACAT CACCGACCGA CTGTCTTCA GCAAGGACAG AGTGCTGATC CAAGTAACCA 2580

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GGCGTCACAT ATTTGGACCA CTCACCTTT CCTTCATCTG GCAGCATTTT ACCAGTCACG	2640
ATACTCATAA AGGTGATTT TCCTTCACCA TTGGCACCGA CCAGGCCGAT ATGTTCTCCC	2700
TTGAGGAGAC GGAAGGACAC ATCTTCAAAA ATTGCACGGT CACCAAAACC GTGACTCAGA	2760
TTTTTAACCT CTAAATACT CATTAAATT CCTTACCTTG TTTTTATGTA ATCGTTTATA	2820
AAGGAGCAA GCCAGATAGC CACCCAAAGT GTGGTCCAC AAATCATCAA TCTCAAAGAC	2880
GCGATTGAAA TCAAAGAAAA AGTCCAAGAT TAATTGCGTA CACTCGATTG CAAGACTCAC	2940
AAGAAAACA AAAAGAAGGA CCTTTTTGT TTTCCGAAA TTTGGAAATA GATAAAGGAG	3000
TTGGAAAATC AGAGGAAAAA ACAAGAACAC ATTGAGGATA TTTGTAAAA AAATCCAACA	3060
TAATTGTCCA ATGTCACTCA CTTCGCCAG TTTCCAGAGA GAATTGAAAG GAGTCAAAG	3120
AAAAACCAGG CGTCCAAGAT GCTGAATACC TGGAGTTCCC ACTCCACGG TAGATTGTT	3180
TTGAGGAGTA AAGCAAAAC AGACAATGCA ATGCTATAG AAAATGACTC CCCAGACCAA	3240
AATATGATTA TAAGTCTTCT TCATCATTAA GGATTTACCG CTGCGACTGC CTTCTGGCGG	3300
TCACGTTCA TTGTGTTAGA GCGCAATTGT CCACAAGCTG CGTCAATATC TGTACCATGC	3360
TCTTGACGAA CCACACAGTT GACCCCTTT TTCTTAAGCG TATCATGAA AGCCAACACG	3420
CACTCTTGG GACTACGGCT ATATTGGTCA TGCTCACTAA CTGGGTTATA AGGAATCAAG	3480
TTTACATAAG ACAATTCTT GATGTTCTT AGCAATTCA GCAATTCCAA GGCTTGTTCT	3540
ACACCGTCGT TGACTTCATT AAGCATGATA TATTCAAAGG TTACACGACG GTTTGTGTC	3600
TCAATGTAGT ATTCAATAGC AGCAAAGAGT TTTCAATCG GAAAGGCACG GTTAATCTTC	3660
ATGATACTTG AACGAAGTTG ATTGTTAGGT GCGTGAAGAG ACACGGCAAG ATTGACCTGA	3720
ACCCCTTCAT CAGCAAAGTC ACGAATTAA TGAGCCAAAC CTGAGGTTGA AACCGTGATG	3780
TGACGAGCAC CGATAGCCAT TCCTTATCA TCATTGATAG TACGAAAGAA ATTCAAGACA	3840
TTGTTGTAAT TATCAAAGGG CTCACCGATT CCCATGACAA CGATATGGCT GATGCGTTCA	3900
TCCTGACCAC GCTCATCAA GTATTTCTGA ACCAGCATGA TTTGCGCTAC GATTCACCG	3960
TTATTGAGGT CACGTTGCTT CTTAATCAA CCAGAGGCAC AGAAGGTACA ACCGATATTA	4020
CAGCCGACCT GAGTGGTCAC ACAGACAGAT AAACCATAGT GTTGACGCAT GAGTACAGTC	4080
TCAATTAAACA TACCGTCGGG CAATTCAAAG AGATATTGTA CTGTACCATC AGCAGACTCT	4140
TGCACAAATAC GTTGTTCAA GGGATTGACC ACAAACTGGT CATTGAGCTT AGCAATCAA	4200
TCCTTGAAA GGTTGGTCAT TTCTTCAT GACTGCACAC GTTTACGGTA GAGCCATTCC	4260
CAGATTGAT CTGCACGGAA TTTCCTTCT CCCTGCTCCA ATACCCATTG CTGCATGGTT	4320
TGATGTACCA AACTATGAAT TGAGGGTTTC ATTTCTTCTC CTTATTCTCT ACTCACTTCT	4380

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GACGAATGAC AAAATGACGT TGTCCTTGT CGTCTTTCTG ACGACGTCTA TTTTCTTAT	4440
CTGCATTCGA CTTCGTTTA GTTGAGTCG GTTTCTTCC TTTTCTAGAA GGTGTTCTT	4500
CTTCCGCTT ACGCATTTC TTGTCAAATG ATGCTCGCTT AGGGGCTTCA TTTTCTAAGA	4560
CAAAATAGGC ACAACCATAA CTACAATACT CTAAAAGGTA GTCTTGTAAA CGACTGATT	4620
TTTCAAGTTT TTCTTCTGTT CGGTCACTCCT TGAAAAAAC TCGTAGGCGA AGCTGTTCGT	4680
TGCTCCAGTC CCCCACGATA TAATCAAAC TGGTTAACAC TTCTGAAAAA CGCTGATTAA	4740
AAGTCGTAC ATCAAAGGCA TCCTTGATAT TTCAACCAA GGAAAAGCT ATCCCTTCCG	4800
TTTCGACCTT GTCCCCGTGT AAATGGAACCT CGGGACCAAGG AAACCTTGTAA TAGTTGTATA	4860
ATTCAGGTGC AATTCTTTT CGCATAGATA TCCTTTTTC ACAGTTACTT AATACTTTAT	4920
TCTACCATAA TTCTAGCAG TTAGCACGTT TCTCATAAAA ATGAAAAAAAG TCTGACGATT	4980
TTGTCAGACC AGAACATTTT AACCTAAAAA GAGAAGAAC ATTCTTCCCT CCAACTATCA	5040
TTATTTAGCA GCTGCGTACA ATTCACTAC TTTATTCCAG TTGATTACTG AAAAGAAAGC	5100
TTTGATGTAG TCAGGACGCA CGTGCAGGTAA TTTCACGTTAG TAAGCATGTT CCCAACGTC	5160
CAAGCCCAAG ATTGGTTTTT TACCTTCTGA GATTGGTGTG TCTTGGTTTG CTGTTGAAGT	5220
CACTTCAAGT TTCCCTTCTT TGTTGACAAC CAACCATGCC CAACCTGAAC CAAACGAGT	5280
TGTTGCTGCT GCAGTGAAGG CTGCTTGAA TTCTTCAAAT GAACCAAATG TTGCATCGAT	5340
TGCTGCTGCC AGTTCTGCTG AAGGAGCTGT TTTCGCGGA GTCATCAATT CCCAGAAAAG	5400
AGCGTGGTTC AAGTGTCCGC CACCATGGTT GATAAGTGT TGACGGATAT CAGCTGGGAT	5460
AGATTCTACA TCAGCAAGCA AGGCTTCAAG GTCTTCACCG ATTTCAAGGTT GTTTTCTAA	5520
AGCTGCATTG GCATTGTTGA CATAAGTTG ATGGTGT	5558

(2) INFORMATION FOR SEQ ID NO: 104:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 6735 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 104:

GGAATTGTAA ATATCATATT GTTTTGACAC CCAAATATCG TCGTCAAATC ATTTATGGCA	60
GATACAAAGC TAGTATCGGA AGAACATACAC GTGACTTATG TGAGCGTAAG GGTGTAATAA	120
TCCATGAAGC GAATGCTGTG TCAGACCATA TTCACATGCT TATCAGTATT CCTCCGAAAC	180

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TTAGTGTTC	GTCCTTATG	GGCTATTAA	AGGCAGAG	CAGTTGATG	ATTTTGATA	240
AGCATGCAA	TTTAAATAC	AAATATGGCA	ATCGAAGTT	TTGGTGTAGA	GGCTATTATG	300
TAGATAACGGT	AGGCCGAAT	CAGAAAGTGA	TACCTGAATA	TATTAGAAT	CAATTACAAG	360
AAGACAGAGT	AGCAGACCAAG	CTCACGTTAT	TCGAGTCAGT	AGATCCGTTT	ACTGGCGAAA	420
TAAATAAGAG	GAAGTAACTA	AGGTGCTTA	GCACCTGCTC	GGGAAAGTGG	TGCGCGAGGA	480
AGCTATTCTG	GTGGGCTTT	GGCCCTGGCC	GGTAGAAGCG	GCTTATAGCC	GCAGAACAAA	540
CCACCAGTTTC	ACACTGGTGG	TTTGATTTA	AAAAACTTGA	TACATAAAA	TAAAAGTCTA	600
TATAAAGGAT	GGTAAATTC	CTGTTGTCCG	ATTGGACAA	TATCCTAAAT	AGTTACAATA	660
TATGGTCTAT	ACTTTTCTT	AGGAGAAAGC	TAGATGTACA	GACGTTGAG	AGATTTGAGG	720
GAGGATCATG	ATCTGCCCA	AAAGCAAATA	GCTACAATAC	TTTCGTTTAC	AAATTCAAGCT	780
TATGCCAAA	TTGAACGGGG	TGAGCATGGC	TTGACGGCTG	ATGTATTGGT	TAAACTCTCA	840
GATTTCTATG	ACGTCAGTAC	AGACTATTAA	TTGGGATTAA	CTGTTTCC	TGATAAAAATT	900
CGCTTTAGAA	ATAATCTCC	TCAATTTCAT	AGAGTTGAA	AATGAGTGAG	ATTTTTTATT	960
TGCCCTTGA	CAACTGAATA	GCCTAAAATG	GTACTTCCT	CATTGTGGA	GCAAATTGAA	1020
ATGGCTCGCC	ATGATAAGAG	CGATTTAAA	ATCATCAATA	AAATAGAGCG	ATACTTTATA	1080
TGCCATGATA	CAAATGATAT	ACAATGATAC	TTCTGACCGT	TCAGCCTGCC	AACGTAAAAG	1140
AGCAGCAAGT	GAATTCTTA	TGATGACTTC	ATCAGTCATG	CCACGTTGAA	TGTGTGAGTT	1200
TGTTAGATAA	ACGCAATTAA	TCCTAAAAG	GTCCCCGAA	CCTTTGAGT	TCTACAGACG	1260
CATCACGTGG	AGTGTGTAAG	CTTGTGCTA	AAACCGTAA	AACCTGGAA	CGAAAGGAAT	1320
AATAGACTTT	CTGCGAAACA	AAAATATAAT	ACAATAAAAC	TATGAATGAT	GAAGCAAGTA	1380
AAACATTGAG	CGATAGCCGT	TTCAAGATCC	TTGTAGGTGT	TCAGCGCACG	ACTTTGAAG	1440
AGATGTTAGC	TGTGTAAAAA	ACAGCTTATC	AACGTAACG	CGCAAAAGGT	GGACGAAAAAA	1500
GCCTTAAAG	CCTAGACGAT	CTCCTTATGG	TAACTATTCA	ATACATGCGA	GAATAGAGCA	1560
CTTATGAACA	AATTGGGGCT	GATTTGGCA	TTCACGAAAG	CAACTTAATC	CGTCGGAGTC	1620
AATGGGTTGA	AGCAACTCTT	ATTCAAAATG	GTTTTACGAT	TTCAAAATTCT	GCCTTAATTC	1680
TGTAAAACA	GTAAAATTG	AAGGATTGTA	AGGTAAGAGT	TTTTTTCTTT	CTGAAAAAAAT	1740
GGTATAATAG	CAATCAAAAC	TAGAAAATAA	AAACGAAATT	GGACAGATT	TGTCTGTATC	1800
CTAGTAGAGT	GGTGATACTA	TGAAGATTAG	TAAGAGGCAC	TTATTAATT	ATTCCATCTT	1860
GATTCCCTAC	TTGCTTTAT	CTATTTGGG	CTTGATTGTG	GTCTATTGCA	CCACCAGTGC	1920
TATTTTAATT	GAAGAAGGCA	AGAGCGCCTT	GCAGTTGGTT	CGAAACCAAG	GAATCTTTG	1980

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GATTGTTAGT TTGATACTGA TTGCCTTAAT TTATAAATTG AGACTAGATT TTTTGAGAAA	2040
TGAGCGACTA ATCATTTCAG TTATATTAAT AGAAATGCTT TTATTGTTCT TGGCTCGTTT	2100
TATTGGTATT TCCGTAAACG GGGCATACGG TTGGATTCG GTTGAGGAA TAACTATTCA	2160
GCCAGCTGAG TACTTAAAAA TCATTATTAT TTGGTATTAA GCTCACCGAT TCTCCAAACA	2220
GCAAGAAGAA ATAGCTACTT ATGATTTCA AGTTTGACT CAAAATCAAT GGCTTCCCCG	2280
TGCTTTAAT GATTGGCGAT TCGTTCTCCT AGTTCTGATT GGAAGTTGG GAATTTCCC	2340
TGATTTAGGA AATGGCGACTA TTTTAGTCCTT GGTTTCCCTG ATTATGTATA CAGTTAGTGG	2400
AATCGCTTAT CGCTGGTTTT CAACCATTCT GGCCTCGTA TCTGCCGCTT CTGCTTTGT	2460
CTTGACCCT ATCAGCCTAA TCGGTGTTGA GACCTTTCA AAAATTCCAG TATTCGGCTA	2520
TGTAGCCAAG CGCTTTAGTG CCTTTTTAA TCCTTTGCC GATCGTGCTG ATGCAGGTCA	2580
CCAGTTAGCT AATTCTTATT TTGCATGGT CAATGGCGGT TGGTTGGTC TAGGTCTTGG	2640
AAACTCGATT GAAAACGAG GTTATTTGCC AGAAGCTCAT ACAGACTTTG TCTTTCTAT	2700
CGTGATTGAA GAATTTGGCT TTGTTGGTC CAGTCTTATT TTAGCTCTCT TGTTTTCAT	2760
GATTTGCGG ATTATCTTGG TCGGTATCCG AGCGGAGAAT CCTTCAATG CCATGGTTGC	2820
ACTCGGTGTC GGAGGGATGA TGTTGGTC GGTTGGTC AATATCGGAG GGATTCGGG	2880
CTTGATTCGA TCTACAGGAG TGACTTTCCC CTTCTTATCC CAGGGTGGAA ATAGCTTCT	2940
AGTCCTATCA GTGGCAGTAG CCTTTGCTT AAATATTGAT GCCAGTGAAA AACCGCCTAA	3000
ATTGTACCGA GAATTGGAAA ATCAACCAAT GAACCTTCTG TTGAAGTAGG ATAAAGAAAG	3060
GATAGTTTAT GTCTCTCAA AAATTAGAAA ATTATAGTAA TAAAAGTGTG TGCAAGAAC	3120
AAAGTCTTGAT TCTAACAGAA TTACTGGAAG ATATTACTAA AAATATGCTT GCCCCAGAGA	3180
CCTTGAAAA AATAATACAG TTGAAAAGAT TATCAACGCA GGAAGATTAT CAAGGTCTAA	3240
ACCGTCTAGT GACTAGCTTA TCAAATGATG AAATGGCTA TATTCACGC TATTTCTCTA	3300
TCTTGCCCTCT TTTGATTAAT ATTCAGAGG ATGTGGATTT AGCTTATGAA ATCAATCATC	3360
AAAATAATAT TGATCAGGAC TATTTAGGTA AAATTATCTAC AACGATTAAA TTGGTAGCAG	3420
AAAAGGAAAA TGCGCTTGAG ATCCTAGAAC ACTTGAATGT TGTCCTGTT TTGACAGCCC	3480
ATCCAACACA AGTCAACGC AAAAGTATGT TGGATTTAAC AAATCATATT CATACTCTT	3540
TGCGTAAATA CCGTGATGTT AAGTTGGGTT TGATCAATAA AGATAAATGG TACAATGATT	3600
TGCGTCGTTA CATCGAAATT ATCATGCAGA CAGACATGAT TCGTGAGAAA AAATTAAAAG	3660
TGACTAACGA AATCACGAAT GCTATGGAAT ATTATAACAG CTCCCTTTTG AAAGCTGTAC	3720

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CTCATTTGAC GACGGAGTAT AAGCGCTTAG CGCAAGCGCA TGGTCTGAAT TTAAAACAGG	3780
CTAAACCAAT CACCATGGGT ATGTGGATAG GTGGTGACCG TGATGGAAAT CCATTTGTTA	3840
CAGCAAAGAC CTTGAACAG TCTGCACTCA CTCAGTGTGA AGTCATCATG AACTACTATG	3900
ATAAAAAGAT TTACCAACTT TATCGTGAAT TTTCTTTC AACTAGCATT GTCAACGTCA	3960
GCAAGCAAGT CAGAGAAATG GCTCGTCAAT CCAAGGATAA CTCGATTTAC CGCGAAAAAG	4020
AGCTTTACCG TCGTGCCTTG TTTGATATTTC AATCAAAAAT TCAGGCAACT AAAACCTATC	4080
TGATTGAGGA TGAAGAAGTT GGGACTCGTT ATGAAACCGC CAATGATTTTC TACAAGGATT	4140
TGATTGCCAT TCGAGATTCT CTACTAGAAA ATAAGGGCGA GTCCCTGATT TCAGGTGATT	4200
TTGTGGAATT ATTGCAGGCA GTAGAGATAT TTGGTTTTA CTTAGCATCA ATTGATATGC	4260
GACAAGACTC TAGCGTCTAT GAAGCCTGTG TGCCAGAACT CTTGAAATCA GCAGGAATTC	4320
ATTCTCGTTA TAGCGAGTTG AGCGAAGAAG AAAAGTGTGA CCTTCTCTTG AAAGAATTAG	4380
AAGAAGATCC CCGAATTCTT TCTGCGACTC ACGCAGAAAA ATCAGAATTAA TTAGCAAAAG	4440
AATTAGCTAT TTTTAAGACG GCTCGTGTG TGAAAGATAA GTTGGGAGAT GATGTCATCC	4500
GTCAGACCAT CATTTCACAT GCAACCAGCC TTTCTGATAT GCTAGAATTAA GCTATTCTGT	4560
TAAAAGAAGT AGGACTGGTG GATA CGGAAA GGGCGCGTGT TCAGATTGTT CCCCTTTTG	4620
AAACAATTGA GAACTTGGAT CATT CAGAGG AAACAATGAG AAAATATCTT TCTCTTAGCC	4680
TTGCCAAAAA ATGGATTGAC TCACGAAATA ACTACCAAGA AATCATGCTT GGCTACTCTC	4740
ACAGTAATAA AGATGGCGGT TACTTGTCA TATGTTGGAC CCTCTACAAAG GCTCAACAAAC	4800
AATTGACTGC TATTGGAGAT GAATTGGCG TTAAGGTTAC CTTCTTCCAT GGTCGTGGTG	4860
GTACTGTGG TCGTGGTGGT GGGCCAACCT ATGAAGCCAT TACATCTCAA CCGCTCAAGT	4920
CTATCAAGGA TCGTATCCGC TTGACGGAGC AGGGTGAAGT AATTGGGAAT AAATACGGTA	4980
ACAAAGACGC CGCTTACTAT AACCTGAAA TGCTAGTATC GGCAGCTATT AACCGTATGA	5040
TTACTCAGAA GAAGAGCGAT ACCAATACCC CAAATCGTTA TGAAACCATT ATGGATCAAG	5100
TAGTGGACCG TAGTTACGAT ATCTACCGTG ATTGGTCTT TGTAATGAG CATTCTATG	5160
ATTATTTCTT CGAGTCAAGT CCAATCAAGG CTATTTCAAG TTTTAATATT GGTTCTCGTC	5220
CAGCCGCTCG TAAGACTATT ACTGAAATCG GTGGTTTGCG TGCCATCCCT TGGGTATTCT	5280
CATGGTCACA GAGTCGTGTT ATGTTCCCTG GATGGTACGG GTTGGGTTCA AGCTTCAGG	5340
AATTATCAA TAAAAATCCA GAGAATATTG CTATCTTACG AGATATGTAC CAAAATTGGC	5400
CTTTCTTCCA ATCGCTTCTT TCAAATGTTG ATATGGTTT GTCAAATCA AATATGAATA	5460
TTGCTTTGA ATATGCTAA CTTTGTGAAG ACAGAGCAAGT TAAGGCCATC TATGAGACTA	5520

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TTTTAAATGA ATGGCAAGTT ACTAAGAACG TTATCTTGGC TATTGAAGGA CATGACGAAC	5580
TCTTAGCTGA CAATCCATAT CTAAAAGCTA GTCTGGATTA CCGTATGCCT TACTTTAATA	5640
TTCTCAACTA TATTCACTTG GAGTTGATTA AACGCCAACG TCGTGGAGAA TTGTCCAGTG	5700
ATCAAGAACG ATTGATTCA ATCACCATCA ACGGAATTGC GACAGGATTG CGTAATTCA	5760
GTTGATAATT TTCAAGAGTG AATGCTAAAA GTGAATATCA AAAAATTCT AATAGACTAT	5820
TGACAAGTAG TTTAAAATG ATATAATTCA ACCATTAGA AAAGTAATCA TACAAACTTT	5880
TTAGAGAGTC TGTGGTAGCT GAAAACAGAT AAGTGGCAAT GATGAAAATT GGGCTGAATG	5940
CTATTTAGAA TTGAAATTAA TAAAAATTG CTAAGCACAC CTTACAGTGC ATCTCGTTAT	6000
TGCGAGACTG AGCGATAGGG AAATTCCTA TAATTGAGGT GGTACCCGC ATCGACGTCC	6060
TCACACAAAGT TTTTGTGTG AGGATTTTT TGATGGAGGT TAGTATGGAA AGAAAACGAT	6120
GGCGTCGCTT GTTTAGATAA GTGAAATATG TTAAAGGAA TAAAAGGAG AAACAGAATG	6180
AAAAATAAAC GTTTAATTGG AATTATTGCT GCATTAGCAG TCTTAGTAGC AGGAAGCTTG	6240
ATTTATTCTT CAATGAATAA ATCAGAAGCT CAGAATAATA AGGATGAGAA GAAAATAACC	6300
AAGATTGGTG TGCTCAATT TGTGAGCCAT CCATCCCTTG ATTTGATTAA TAAAGGGATC	6360
CAAGATGGAC TTGCGAGAAGA AGGATATAA GATGATCAAG TTAAAATTGA TTTTATGAAC	6420
TCAGAAGGTG ACCAAAGTAA GGTTGCGACA ATGAGTAAAC ATTGGTTGC AAATGGGAAT	6480
GACCTTGTGG TTGGTATCGC AACACCAGCA GCCCAAGGGT TGGCTAGTGC AACAAAAGAC	6540
CTACCGGTTA TCATGGCCGC TATTACAGAC CCAATTGGTG CTAACCTGGT TAAAGATTTG	6600
AAAAAACCAAG GTGGCAACGT TACAGGGTA TCTGACCACA ATCCAGCTCA ACAACAAGTT	6660
GAACTCATCA AGGCTCTGAC ACCGAATGTG AAAACAATCG GAGCTCTTA CTCAAGTAGC	6720
GAAGACAATT CAAAA	6735

(2) INFORMATION FOR SEQ ID NO: 105:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 6516 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 105:

CTAGAGGATC CCAGCAGGTA AATTGGCTTC AGCTGGCAA AAAGTTGCC TCGTTGAACG	60
CAGCAAGGCT ATGTACGGTG GAACTTGTAT CAACATTGGT TGTATCCAA CTAAACCTT	120

778	
GCTAGTTGCT GCTGAAAAGG ACTTGTCTTT TGAGAAGTC ATTGCTACTA AAAACACGAT	180
CACTGGTCGC CTCACCGTA AAAACTATGC GACTGTTGCT GGTACAGGCG TAGATATCTT	240
TGATGCCGAA GCTCACCTCC TTTCAAATAA AGTCATCGAA ATCCAAGCTG GTGATGAAAA	300
GAAAGAACTG ACTGCTGAAA CAATCGTCAT CAACACTGGT GCTGTTCAA ACGTCTTGCC	360
AATCCCTGGA CTTGCTACAA GCACAAACAT CTTTGACTCA ACAGGTATCC AAAGCTTGGA	420
CAAATTACCT GAAAAACTTG GAATCCTTGG TGCGGAAAT ATCGGTCTTG AATTTGCCGG	480
CCTTTACAAC AAACCTGGAA GCAAGGTCAC AGTCCTAGAT GCCTTGGATA CATTCCCTACC	540
TCGTGCAGAA CCTTCCCATCG CAGCTCTTGC TAAACAATAC ATGGAAGAAC ATGGCATTGA	600
ATTGCTTCAA AATATCCATA CTACTGAAAT CAAAAACGAT GGTGACCAAG TGCTTGTGCGT	660
AACTGAAGAC GAAACCTTACC GTTTGACGC CCTCTCTAC GCAACTGGAC GCAAACCCAA	720
TGTTAGAACCA CTTCAACTTG AAAATACAGA TATTGAACTA ACTGAACGTG GTGCTATTAA	780
AGTAGACAAA CACTGTCAA CAAACGTTCC TGGTGTCTTT GCAGTTGGAG ATGTCAACGG	840
TGGCCTTCAA TTTACTTACA TTTCACTTGA TGACTTCCGT GTTGTAAACA GCTACCTTGC	900
TGGAGATGGC AGCTATACAC TTGAAGACCG TCTCAATGTG CCAAATACTA TGTTCATCAC	960
ACCTGCACTT TCACAAGTTG GTTTGACTGA AAGCCAAGCA GCTGATTGA AACTTCCATA	1020
CGCTGTTAACG GAAATCCCCG TTGCAGCAAT GCCTCGTGGT CACGTTAAATG GAGACCTTCG	1080
CGGTGCCCTTC AAAGCTGTTG TCAATACTGA AACAAAAGAA ATTCTTGGAG CAAGCATCTT	1140
CTCAGAAGGT TCTCAAGAAA TCATCAACAT CATCACTGTT GCTATGGACA ACAAGATTCC	1200
TTACACTTAC TTCACAAAAC AAATCTTCAC TCACCCAACC TTGGCTGAGA ACTTGAAATGA	1260
CTTGTTGCG ATTTAAGTTG AGATTTAACG GTATCGAACA GCCCTCTTTG GGCTGTTTT	1320
ACTTCTGCGG AATCTCAAAT CTGTCCTTCT CCTCTTTAT GATATAATAG AAACATGAAC	1380
TTAAAAACTA CTTTGGGCCT TCTTGTGGG CGTTCTTCCC ACTTCGTTTT AAGCCGTCTT	1440
GGACGTGGAA GTACGCTCCC AGGGAAAGTC GCCCTTCAAT TTGATAAAAGA TATTTTACAA	1500
AACCTAGCTA AGAACTACGA GATTGTCGTT GTCACTGGAA CAAATGGAA AACCCGTACA	1560
ACTGCCCTCA CTGTCGGCAT TTTAAAGAG GTTTATGGTC AAGTTCTAAC CAACCCAAGC	1620
GGTGCCAAACA TGATTACAGG GATTGCAACA ACCTTCTAA CAGCCAAATC TTCTAAAATC	1680
GGGAAAAATA TTGCGCTCCT CGAAATTGAC GAAGGCCAGTC TATCTCGTAT CTGTGACTAT	1740
ATCCAGCTA GTCTTTTGT CATTACTAAT ATCTTCCGTG ACCAGATGGA CCGTTTCGGT	1800
GAAATCTATA CTACCTATAA CATGATATTG GATGCCATTG GGAAAGTTCC AACTGCTACT	1860
GTTCTCCTTA ACGGAGACAG TCCACTTTTC TACAAGCCAA CTATTCCAAA CCCTATAGAG	1920

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TATTTTGGTT TTGACTTGG AAGGGACCA GCCCAACTGG CTCACTACAA TACCGAAGGG	1980
ATTCTCTGTC CTGACTGCCA AGGCATCCTC AAATATGAGC ATAATACCTA TGCAAACCTG	2040
GGTGCTATA TCTGTGAAGG TTGTGGATGT AAACGTCCCTG ATCTCGACTA TCCTTGACA	2100
AAACTGGTT AGTTGACCA CAATCGCTCT CGCTTGTCA TAGACGGCCA AGAATACGGT	2160
ATCCAAATCG CGGGCTCTA TAATATCTA AACGCCCTAG CTGCTGTGGC CATGCCCGT	2220
TTCCCTAGGTG CCGATTCGCA ACTCATCAA CAGGGATTG ACAAGAGCCG TGCTGTCTT	2280
GGACGCCAAG AACCTTTCA TATCGGTGAC AAGGAATGTA CCCTGTCTT GATTAAAAAT	2340
CCAGTCGGTG CAACCCAAGC TATCGAAATG ATCAAACCTAG CACCTTATCC ATTTAGCCTA	2400
TCTGTCCCTCC TTAATGCCA CTATGCAGAT GGAATTGACA CTAGCTGGAT CTGGGATGCA	2460
GACTTTGAAAC AAATCACTGA CATGGACATT CCTGAAATCA ACGCTGGCGG TGTCGTCAT	2520
TCTGAAATCG CTCGTCGCCT CCGAGTGACT GGCTATCCAG CTGAGAAAAT CACTGAAACG	2580
AGTAATCTGG AGCAAGTTCT CAAGACCATT GAGAACATCAAG ACTGCAAGCA TGCCATATT	2640
CTGGCAACTT ATACTGCCAT GCTGGAATTG CGTGAACCTGC TGGCTAGTCG TCAGATTGTT	2700
AGAAAGGAGA TGAACATAATG GTTTATACTT CACTTTCCCTC AAAAGATGGC AATTACCCCT	2760
ATCAGCTCAA CATTGCCAC CTCTACGGAA ATCTCATGAA TACTACGGGG ACAATGGAAA	2820
CATCCCTCATG CTCAAGTATG TGGCTGAAAA ACTGGGAGCC CATGTGACCG TTGACATCGT	2880
TTCTCTCCAT GATGACTTTG ATGAAAATCA CTACGACATC GCCTTTTCG GTGGTGGTCA	2940
AGACTTTGAA CAAAGTATCA TTGCAGACGA CCTACCTGCT AAAAAGAGA GCATTGACAA	3000
CTACATCCAA AACGACGGTG TAGTTCTGGC TATCTGCGGT GGTTCCAAC TATTGGGTCA	3060
ATATTATGTT GAAGCTTCAG GAAAACGTAT CGAAGGGCTA GGGGTCAATGG GACACTACAC	3120
GCTCAACCAG ACCAATAACC GTTTTATCGG TGACATCAAG ATTACAAATG AAGATTTCGA	3180
TGAAACCTAC TATGGATTG AAAATCACCA AGTCGTACC TTCTCTCTG ATGACCAAAA	3240
ACCGCTGGGA CAGGTTGTCT ATGGAAATGG AAACAACGAA GAAAAGGTCG GTGAAGGGGT	3300
TCATTATAAG AATCTCTTGT GTTCTACTT CCACGGGCCT ATCCTCTCTC GTAATGCCAA	3360
TCTGGCTTAT CGCCTAGTTA CTACTGCCCT CAAGAAGAAA TATGGTCAGG ACATCCAAC	3420
CCCTGCCTAT GAGGACATTC TCAGCCAAGA AATCGCTGAA GAGTACAGTG ACGTCAAAG	3480
CAAGGCTGAC TTTTCTTAA CAAAGGAAA TGATATCAAA GAACTCCGTT ATCTGTGCGG	3540
AGTTTTTGT CTTTCTTTT ACCCTCTCC CTTGCATTTT CTCTCATTTT TTGCCAAAAT	3600
AGAGGGGTAG AAAGAAGGTA GCATATGTCT AAATTACAAC AAATCCTAAC ATATCTTGAA	3660

780

TCAGAAAAAC TAGACGTCGC TGTGCTATCT GACCCCGTCA CAATCAATTAA CCTCACTGGT	3720
TTTTACAGTG ATCCCCATGA ACGCCAAATG TTCCCTCTTG TCCTAGCAGA TCAGGAACCT	3780
CTCCCTCTTG TCCCAGCTCT TGAAGTAGAA CGTGCAAGTA GCACCGTTTC CTTCCCAGTA	3840
GTGGGCTATG TCGATTCTGA AAATCCATGG CAAAAAAATCA AACATGCTCT TCCACAACCTT	3900
GACTTCAAAC GTGTCGCTGT TGAGTTTGAC AATCTCATCT TGACCAAATA CCATGGTTTG	3960
AAAACAGTTT TTGAGACTGC TGAGTTTGAC AACCTCACTC CTCGTATCCA ACGCATGCGC	4020
CTCATCAAAT CAGCTGATGA AGTGCACAAA ATGATGGTTG CAGGTCTTTA TGCTGACAAG	4080
GCTGTTCATG TTGGTTTGA CAATATTCT CTTGATAAGA CTGAGACAGA TATCATCGCA	4140
CAAATCGACT TTGCCATGAA ACGTGAAGGT TATGAAATGA GCTTTGATAC CATGGTCTTG	4200
ACTGGTGATA ATGCTCGAA TCCACACGGC ATTCAGCAG CTAATAAGGT TGAAAATGAT	4260
GCTCTTCCTCC TCTTTGACCT GGGTGTCTG GTCAATGGCT ATGGCTCAGA TATGACTCGT	4320
ACAGTCGCTG TCGGCAAACC AGACCAATTIC AAGAAAGATA TTTACAACCTT GACTCTTGAA	4380
GCCCAACAAG CTGCTCTTGA CTTTATCAAG CCAGGTGTGA CTGCTCATGA AGTGGACCGC	4440
GCTGCCGCTG AGGTCAATCGA AAAAGCTGGT TATGGTGAGT ACTTCAACCA CCGTCTCGGG	4500
CATGGTATCG GTATGGATGT CCATGAATTIC CCATCTATCA TGGAAGGAAA CGACATGGTC	4560
ATCGAAGAACG GCATGTGCTT CTCTGTTGAA CCAGGTATCT ATATCCCTGG TAAAGTCGGT	4620
GTTCGTATTG AAGACTGCGG TGTGTTTACG AAGGATGGCT TCAACCTCTT TACAAGCACC	4680
AGCAAAGATT TGCTTTATTT TGATTAACAT ATATAGCCCC TATGCTTCC TTTCAAAATA	4740
TCTAGGGCT ATTTCATTGT CATTTCCTG CTATTATGCT AAAGAAATTG GCTGCAATAA	4800
TCTAACCCCTA ATGCTCTGGA ATGATAACGA GGGTGCTCTC CGCTTTATC AAAGACAAGG	4860
GATGAAACCC CAAGAAACAA CAATGGAAAT GATAATTGAT TAAGAAGTCA TCTATCAAAA	4920
GATGTTAGAA AAAGTTCAAT TTCACTAGAA AATGAGGAAA ATCTCCCCAC AATAAAACGC	4980
ATAGTATCAG GTATTGTGTA CTGACCCCAA ACAGTTAGAC AATTAATTAA TCCGAAGGAT	5040
TTAGTTCTGT ACTGCACAGG ACTAAGTCCT TTTAGTTTA CCTTAATTG TTTGTTGTTG	5100
TAGTAATCAA TATAGTCTAT AATGACTTGT TCCAATTGGT TAAAGTGTATT AAATGTTTC	5160
TCATAGGCCAT AAAACATTTC GGATTTAAA ATGCCAAAGA AAGATTCCAT CATACCGTTG	5220
TCTTGGCTGT TTCCCTTGCG TGACATAGAT GCTTGAATTG CTTTATTCTC TAGGAACCGA	5280
TGATAAGAAT CGTGTGGTA TTGCCAGCCT TGGTCACTAT GGAGAATCGT ATTCTCGTAG	5340
TGCTTCTCTT TGAATGCCCTG TTCCAACATT GTTTGTACTT ATTCTAAATT AGGCGAACAA	5400
GAAAGATTAA AAGCAATAAT TTGGCTGTTA AAGCCATCTA AACTGGTGA TAAGTAAAGC	5460

781

TTTGAGTAC TTGCTGGAAT GGCAAATTCA GTCACATCG TGTAGCACTT TTCCATTGTT	5520
TTAGAGCCTT CAAATTGGGC TTGAATGAGA TTCTCTGCCT TCTTACCAAC GTCTCCTTTA	5580
TGAGAAGAAT ATTTTCGTTT CTTCGCATT TTAGCTTGTA AATTGAGTAC TTTCATCAAG	5640
CCTTGAACTC TTTTATGATT TACAGATAA CCACGATTC TTAGTTCTAA ATGAACCCGG	5700
CGATAAGCAT AATTCCCTT GTGTCGATA AAGATGGATT GAATTTCACT TTTAAGCTCT	5760
TGGTCTTAT CTGTTTGTC TAGCTGTTTC AAGTGATAGT AGTAGGTCCA ACGAGCTAGT	5820
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GTCTTCTTC TTTCTCTTTT TCCTCCTTCA ATCGGAGTTC TCTTAACCTT TTTAGGATGG	5940
CATTCTCCGC TCTCAGGTAC TCTCCCTCTT GTTTCTCAA CAATAGTATA CCCGTTTTC	6000
CTGTATTGTG CTAGCCAGTT AAGAAGTATC GTACGACTTG GGAGACCGTA TTCAAGAGAA	6060
ACTCTATCTT TAGCCAGCC TTCATGTCAG ACTTTATTAA CCCCATTAT TCACCCCCAA	6120
TCTAAAAACC ATCCAGAATC CTTGCCTTAG CTTAGATCCT GGATGGTTTC TTTTTTCACC	6180
CAATGGGTGT TTTTACTAG ACAAAAAGA GTTTCCCTT TATGGTATAA GTGTAGAAAA	6240
AAACACAAAA AGAAAGGAAA CTCACATGAA CAGTTTACCA AATCATCACT TCCAAACAA	6300
GTCTTTTAC CAACTATCTT TCGATGGAGG TCATTTAACC CAGTATGGTG GTCTTATCTT	6360
TTTCAGGAA CTTTTTCCC AGTGAAACT AAAAGAGCGG ATTTCTAAGT ATTTAGTAAC	6420
GAATGAAAGCGCTACT GTCGTTATTC GGATTCAGAT ATCCwTGTCAG ATTCCTCTT	6480
TCAACTGTTA ACAGGTTATG GAACGGAATA TGCTTG	6516

(2) INFORMATION FOR SEQ ID NO: 106:

- (i) SEQUENCE CHARACTERISTICS:
- (A) LENGTH: 14654 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 106:

TTTTCAACCC ATATCGTGGC TCCTGAATAC TACTTACTGA CAACTATGCT ATCAGAGACT	60
TCTCTACTTG TTTTCTATAT CATTTCATC CATAGAAAAC AACTCATCCA CTTGGGACAT	120
ATCTTTAGCT ATACTGTTCG ATACTCTCTC TTTTCACTTT CCTTTGTAGC AATTATTTTC	180
CTGATTAATT TCGTGTATCC TGTAGATATG GTCATTAATT TGCCATTTT GATTAATACT	240
GGTTTGATTG TCTTGCTATC AGCTATCTCT TATATTAGTC TACTTGCTTT CACAAAAGAT	300

782	
ACCATTTCT ATGAATTTT AAACCATGTC CTAGCCTTAA AAAATAAATT TAAAAAAATCA	360
TAGGAGTTA AAATGAAACA ACTAACCGTT GAAGATGCCA AACAAATTGA ATTAGAAATT	420
TTGGATTATA TTGATACTCT CTGTAAAAG CACAATATCA ACTATATTAT TAACTACGGT	480
ACTCTGATTG GGGCGGTTCG ACATGAGGGC TTTATCCCTT GGGACGACGA TATTGATCTG	540
TCCATGCCTA GAGAAGACTA CCAACGATT ATTAAACATT TTCAAAAGGA AAAAGCAAG	600
TATAAGCTCC TATCCTTAGA AACTGATAAG AACTACTTTA ACAACTTTAT CAAGATAACC	660
GACAGTACGA CTAAAATTAT TGATACTCGA AATACAAAAA CCTATGAGTC TGGTATCTTT	720
ATCGATATTT TCCCTATAGA TCGCTTGAT GATCCTAAGG TCATTGATAC TTGTTATAAA	780
CTGGAAAGCT TCAAACGTCT GTCTTTCAAGT AAACATAAAA ATATTGTCTA TAAGGATAGC	840
CTTTTAAAG ATTGGATACG AACAGCCTTC TGGTTACTCC TTGACCGGT TTCTCCTCGT	900
TATTTGCAA ATAAAATCGA GAAAGAAATT CAAAAATATA GTGCGAAAA TGGGCAATAT	960
ATGGCTTTA TCCCTTCAAA ATTTAAGGAA AAGGAAGTCT TCCCAAGTGG TACCTTTGAT	1020
AAAACAATCG ATTTACCCCTT TGAGAATTAA AGCCTTCCTG CACCTGAAAA ATTTGATACT	1080
ATTTTGACAC AATTTTATGG AGATTATATG ACCCTACAC CAGAAGAAAA ACGCTTCTAC	1140
AGTCATGAAT TTCACGCTTA TAAATTGGAG GATTAGGATG CAATATTAG AAAAAAAAGA	1200
AATTAAGAA ATTCAACTAG CCCTGCTGGA CTATATTGAT GAGACTTGTAA AGAAAACATGA	1260
TATTCCTTAT TTTCTCAGTT ATGGAACCAT GCTTGGAGCC ATCCGCCACA AAGGTATGAT	1320
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GATTATTGAA GAAGAAAATC ACCCTCGCTA CAAGGTTCTT TCCTACGATA CATCTTCTTG	1440
GTACTTCCAT AATTCGCAT CGATTTGGAA CACTTCTACT GTTATAGAAG ACCATGTTAA	1500
GTACAAGCGT CATGATACCA GCCTTTTCAT CGATGTCCTTC CCAATTGATC GATTACAGA	1560
CTTGAGCATT GTCGACAAGA GCTATAAGTA TGTGGCTCTT CGTCAACTAG CTTATATCAA	1620
AAAATCACGA GCAGTTCAAG GTGATAGCAA ACTAAAGAT TTTCTTAGAT TATGTAGCTG	1680
GTACGCTCTC CGATTTGTCA ATCCTCGCTA CTTTTACAAG AAAATTGATC AACTAGTCAA	1740
AAATGCTGTA ACCAACACTC CTCAATATGA AGGAGGAGTT GGGATCGGT AGGAAGGGAT	1800
GAAAGAAATC TTCCCAGTTG ATACCTTAA AGAACTGATT TTAACTGAGT TTGAGGGCCG	1860
TATGTTGCCT GTTCCCAAAA AATATGACCA ATTTTAACC CAGATGTATG GCGATTATAT	1920
GACACCACCA TCAAAAGAAA TGCAAGAGTG GTATAGTCAT AGCATTAAAG CTTATCGCAA	1980
AAACTGATTG AGGGGGATTA TACAAACTAC TAAGATAGAG GTTATTCAA AACATAATT	2040
TAGTAGAAAA TGAAATACAT ATTCCCACAA TAAAACGCAT CATATCAAGG TTTTTGAAAA	2100

783

ACCTTGATAT GATGCGTTT ATAATTTAA AGACTTTTT CTATAGTAGA TTGAAATAAG	2160
ATGCGAACAA ATCAATTAGA AAATTCAAAT TAATTTATAG AAATATTTA GTATTCCTGT	2220
GTACTGTTCT AAATTCAAGTC TGCTATATCT TATTTTCTA TTTAAATCGC TTCTGTAACA	2280
AAGCTACGAC TTTCAAGTAC CTTAACGCATG GCATTTAGCTG TATCTAGCGC TGTGAAGAGG	2340
GGCACCCCGT GTTCAATGGC TGAACGACGA ATTTGCTCAC CATCTTCGTC AGCAGTCGT	2400
TTTGTCCCTA CTGTGTTAAT GATAGCTTGA ATTCTTCCTT TGCGTACAAA ACTTGGGATA	2460
TCCTTATCGT CATCACCAAT CTTACCAACA GGTTGGGCTT GCAAGCCATG ACTAGCAAAG	2520
AAGGCTGCTG TCCCTCTGT CGCAAGGATT CCATAACCAA TGTTTGGAA ACGACGAGCC	2580
AAGTTCAAGG CTTCTCTTT GGCAATCATCA GCGATGGTAA AGACGACATT ACCAAAAGTT	2640
GGCAAGTGTAA GATAAGAAGC TTCAAAGGCT TTATAGAGAG CTTTTCCAA AGTAGCATCA	2700
GAACCCATAA CTTCACCTGT TGACTTCATT TCAGGACCGA GCAAGCTGTC TACCTTAGCT	2760
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CCATTTGGT AGCCAAGTTC TGATAAACATT TGACCAAGAA TGAGTTGGT CGCTACTTGA	2880
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ACCTCAATAA CGTAGACTTT TTCATCCTTG ATAACAAACT GGATGTTCAT CATTCCAAGG	3000
CAGTGAAGAC CGATTGCTAA GCGTTTGGTG TAGTCTGCGA TGGTCTCCTG AACCTTTGC	3060
GACAAGGTTT GTGGTGGTA AACAGCCATT GAGTCACCTG AGTGGACACC AGCACGTTG	3120
ATATGCTCCA TGATACCAGG AATGAGTACA TTTTTACCAT CTGAAATGGC ATCAACTTCG	3180
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GCAGTTCGCA TGTAAGAACG AAGGTCTTCT TCGTTTCAA CGATTCCTG GGCACGTC	3300
CCAATACAT AAGATGGCG GACAAGAACT GGGAAAGCCAA TCTTGCAGAC TGCAAGAGCT	3360
GCTTCTCTT CATTGGTAGC CGTTTGTCTT GGTGGCTGTG GAATATCCAA TTCTTTGAGA	3420
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TGAACGATAA CTCCCTTGG TTGTTCCAAG TCAATGACGT TCATAAACATC TTCGAATGTC	3600
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GGTAGTCAA ACTCAACCCC TTGACCGATA CGGATTGGAC CTGAACCTAG GACAAGTACA	3780
GATTCTTAT CAGATCTGAT AGATTCAATT TCCCAACCAT AGTTGAATA GAAATATGGC	3840

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GTTTCGGAGT CGAACTCTGC CGCACAAAGTG TCTACCATCT TATAAACTGG AACAAATCTTG	3900
TTTTCCAAGC GAAGTGGCG AACTTTATCA TCAGTCGTT CCCAGAGTTC AGCAATCTTA	3960
CGGTCTGAAA AACCATTAAG TTTGCTGTT TTCAAAACTT CTAATCTTG TGATGAGCA	4020
CCCAATTCTT GCTCAATTTC AAAGATATGC AAGAGTTAT CAAGATAGAA GATATCAATT	4080
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AAGAGACGGT CATCTGGGC TTTGACAACC TTTCAATCA AGGCATCATC AGAAACTGCT	4200
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ACGTAGTCAA GGGCTGGTTC AAACATGGCA TAGGTTGAAC CTGTAACTGG GTTATAACC	4440
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CGAATAATGC TCAAGCTCGC ATCACGAAGC ATTTGGTTTT CATACTGTA CATGGTTTGCA	4680
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GCCTGAGTCC CAGCGTAGTC AAACTCAGCA GCCTGACCAA TAATAATCGG ACCAGAACCA	5400
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CTATTGCTCG CCTCTCAGGG CGACATTAAA TAAGATACAA AGGACGAATA GAAAGCGATT	5640

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GAATTTAGG AAATCAAGGA AGGATTGACA ATCCAAGTTG GTTTCTCTAC ATTCTGAGCT	5700
TTCCGTCCGT GTTCAGTTAC ATAATTCTC CGACGAGCTT TTACTCGTTC TTAGTTGAT	5760
TGTTTAAAAA CTTCCATCAT CTCGATAAAC TCGTCAAATA GGTAGCTAGC GTCGTGTGGC	5820
CCAGGAGCTG CATCTGGGTG GTATTGAACA GAGAAAGCAG GTGGTATCT GTGGCGCACA	5880
CCTTCCACTG ACTTGTCATT GATTCTTCG TGGAATAAA TCAAGTGCTC TGCAAATCC	5940
TCGCGGCTGA CTGCATAACC ATGGTCTGG CTGGTGAAGT CTACTCGTCC TGTTGCGATT	6000
TCACGTACCG CATGGTGAA TCCACGGTGG CCAAACCTCA TCTTATAGGT CTTAGCCCCG	6060
TTTGCCATG CAAAGAGTTG GTGTCCTACAAACAGATTGAAATTTCCCTTGAT	6120
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AACATAACTC CGTCAGGATT GAGATGGAGA ATTTCTTCAG CCGTTGTGCA ATAAGGAACA	6240
ACTGTCACGT TACAGTTGCG TTTAGAAAATG TCACGTAGGA TTGAGTGCTT GAGACCAAAG	6300
TCCACTAGCA CCACGCTAA ACCAACTCCT GGAGCTGGAT AAGACGTTTT AGTAGAAACC	6360
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ATAGTCTGCA AACTCTTGAC TTCTCCATTC CTCAGGTCCA GCGAAAAAGA GTTCAGCTCC	7260
CAAGCGTTTC AAAATCTGCA TATTGGATTT GGCAACGCGT GAGTGGTCCA AGTCACCTGC	7320
AATAGCAACT TTAAGACCCCT CAAAGTGGCC AAATTCCCTCA TAAATGGTCA TCAAATCAAG	7380

786	
CAAGCTCTGG CTAGGGTGTG GGCCCGAACCC ATCTCCACCA TTGATGATGG AAGTCGTAAT	7440
CGTTGGACTA GCAATCAATT CTCTATAGTA GTCGACCTCT GGATGGCGAA TCACACAGAC	7500
ATCCACTCCT AAAGCAGACA GAGTCAAAAT GGTGTCTAA AGTGTCTCAC CCTTATTAAC	7560
CGAGCTAGTC TTCACATCAA AGTCAAGTCG TTCCAATCCA AGTTTAATCT CTGCGACTTC	7620
AAAGGACTTA TGTGTCCGTG TAGAACCTC AAAGAAGAGA TTGGAAACAA TCGGATGGTC	7680
TTCATAGGGA AGCTGGGCTC CATTTTAAA CTCATTCT CGCTTGATCA ATTTCATTAC	7740
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AAGTTCTGTC ATCTCTACGA TGATTTCTTC AGAACGACTG GTTGGGATAT TTTTCCAAC	7920
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TGATAACATG ACCGCAACCA GGAGTTCAAATGATTGGTA AAATCAAGAC TAGGCTTGGC	9120
ATCTGGGAAG AGGGCAATGA TTTCTCTAG CACCTTTCGT GCTCGTTTT TTGACAAGAC	9180

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CATTATTCAT CTCCGTCAA TAGTCCTTGT AAGGCCAGCAA AAGGACTGTT TTCTTCCTTTC	9240
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ATAAAATCCTT GACCAGCTTC TTCTTCAGCC GTCAAGACCT TGATAGGAAT GTTTAGCAGG	9360
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GCAAGGATAT CTTTACATC TAAAATTCT TGATTACGTG CACGCAGGTC ATCAACTAAA	9660
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AGGGGGCGAA GCTTGAAAAA TTCAAGCTTAT CCATGAGTTA GCCAAACCCCT TTGAGATTCT	10920

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ATTTTTAGAT GAACCTCAA ATGACCTAGA CCTTGAGACA GTTGATTGGC TAAAAGCCA	10980
GATTCAAAAG ACCAGGCAAA CGGTTATTTT CATTTCCTCAT GATGAAGACT TTCTTTCTGA	11040
AACGGCAGAC ACTATTGTTC ACTTGCGACT GGTCAAACAC CGTAAAGAAG CGGAAACGCT	11100
AGTAGAGCAT TTAGACTATG ATAGCTATAG TGAGCAGAGA AAGGCTAATT TTGCCAACAA	11160
AAGTCAGCAA GCTGCTAACCA ACCAAAGAGC CTACGATAAA ACCATGGAAA AACATCGGAG	11220
AGTTAAGCAA AATGTAGAAA CTGCGCTTCG AGCTACCCAA GATAGTACTG CCGGTCGCCT	11280
ATTGGCTAAA AAGATGAAAA CTGTCCTCTC ACAAGAAAAA CGCTACGAAA AGGCAGCTCA	11340
GTCCATGACT CAAAAGCCAC TTGAAGAGGA ACAAAATCCAA CTTTTCTTTT CAGACATCCA	11400
ACCATTACCA GCTTCTAAAG TCTTAGTCCA ACTGGAAAAA GAAAATTGT CCATTGACGA	11460
CCGAGTTTG GTTCAAAAAC TACAACTAAC TGTCCGTGGC CAAGAAAAA TCGGTATTAT	11520
CGGGCCAAAT GGTGTTGGGA AATCAACTCT GTTAGCCAAG TTACAGAGAC TTCTGAATGA	11580
TAAAAGAGAG ATTTCACTTG GTTTTATGCC ACAAGATTAC CACAAAAAAAC TGCAATTGGA	11640
TTTATCCCCA ATAGCCTATC TCAGTAAAC TGGGGAAAAA GAGGAAC TAC AGAAAATCCA	11700
ATCTCACCTA GCTAGTCTCA ATTTCACTTA TCCAGAAAATG CAGCATCAAA TTCGCTCCTT	11760
ATCTGGCGGA CAACAGGGAA AACTCCTGCT TTTGGATTAA GTCTGCGCA AACCAAACCTT	11820
TCTCCTGCTG GATGAACCCCA CACGAAACTT TTCTCCCACT TCTCAACCCC AAATCAGAAA	11880
ACTCTTGCT ACCTATCCAG GCGGTCTCAT CACTGTTCG CATGACCGTC GTTTCTTAAA	11940
AGAAGTCTGC TCGATCATCT ATCGCATGAC AGAACACGGT TTGAAGCTAG TTAATTAGA	12000
AGATTATAA ATTTGCAACA TAGCAAAAT CCAGAGACGA CCTCTGGATT CTTTTACATC	12060
TGTTTAAAC GTTCAATCCG TTCTGAGATA GGTGGGTGGG TATAAAAGAG TTTTTGGAAC	12120
CCCCCACCTT TCTTAGGATC ATTGATATAA AGGGCACTGC TAGCATCATC GACGTGGCGA	12180
CTCATAGTT TGCTATTGTC CAACTTATCT AGGGCATTAA TCATTCCCTG GGGATTGCCA	12240
GTCAGCTCGA CACTAGATGC ATCTGCCAGA AATTCCCTCT GACGAGAAAT AGCGAGCTGA	12300
ACCAAGGTTG CAGCGAGAGG TGCCAGTACA ATAGCTAGTA GGGAAACAC TAGCATAATG	12360
ATTTCAAGAC CATTTCATC TCGGTCTCATCA TCACCTCGTC TGCGACCTGC TCCACCCAC	12420
CACATCATAC GACCTGCCAT ACTAGAAAGC ATGGTGATAG CACTAGCAAG GGCAACTGCA	12480
ATAGTCGAAA TACGGATATC ATAATTACGA ATATGACTGA CTTCATGTCC CATAACAGCT	12540
TCTAGTTCTT CACGATTCTAT GATAGCTAGT AGACCTGAAG TCGCAGCAAC AGCCGCATT	12600
TGAGGATTAG AACCTGTCGC AAAGGCATT AAGGCTGGAT CATCAATGAT GAAAACACGG	12660
GGCATAGGAA TCTGAGCGAC CAGAGCCATA TCTTCCACTA CATGGTAGAG GTCTGGTGC	12720

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GTTTGCTCAT CCACCTCACG CGCTCCATTC ATGGACATGA CAATCTCTGT CGATTGAAAA	12780
ATCATAGACA AAGCGTAGAT AAAGCCGATA ATCAGTGCAA TAACCAAACC ACCAAGTCCA	12840
GATCTTATAA AGAGATAACC AACCGCATAA CCAACAAGAG CTAAGAGTAG GAAAAATACC	12900
AGCAACAAAA TCCAGGTTT TCGTTTATTC CTTGCAATTG GATCAAACAA CATCTTAGTC	12960
ACCTAAACCG CTAAAATCAA CTTTAGGAAC CGACTTTCC TCTTCAGGTG TTTGAAGGAA	13020
ATCTGCCGCT TTAAATCCAA ACATTCCAGC GATAATATTG CTCGGAAAG TTTCTAATTT	13080
TACATTGTAG TTGCTGACAA CACTGTTATA GAGTTGACGA GAGTAAGAAA TTTTATTTTC	13140
TGTGTTGTC AACTCCCTT GCAATTAAAC AAAGTTAGCA CTAGCTTC AATCTGGATA	13200
GCTTCTGCA ACTGCAAAAA TACCTGAAAC CTGACGAGTG AGGGCATCAC TGGCTTCAT	13260
AGCTTCTGCT GGTGAAGTCG CTGCCGCCAC TTGGTTACGT AGTTCTGCCA CCTTTCAAG	13320
GGTAGAACCT TCATATTTGG CATAACCTTT TACAGTCTCA ATCAAGTTTG GCAAGAGGTC	13380
ATTGCGACGT TTCAACTGAA CATCAATCTG ACTCCAAGCC TCCCTGGTT GCATACGATT	13440
TTTAACCAA CGTTTATAGC TAACAATCAC AAAATAACA ATAAGAGCGA TAACTCCAAG	13500
AATAATCCAA GTCATAATAT AAGTCCTTTC TGCTTTAGA TTAGTACCAAG TATATCAAAT	13560
TTTCTATGAT TGTGGAAAAA TAAGATGATA CTAAAGAAGG AAATAACTAT GAAACCAAAA	13620
ACATTTTACA ACTTGCTTGC CGAGCAGAAT CTTCCACTTT CGGACCAGCA AAAAGAACAA	13680
TTTGAACGTT ATTTTGAGCT CTGGTTCGAG TGGATGAGA AGATTAATTG GACGGCGATT	13740
ACGGACAAGG AAGAAGTTA TCTCAAACAT TTTTACGATT CGATTGCACC CATTCTCAA	13800
GGTTGATTC CCAATGAAAC TATCAAACCTT CTTGATATCG GGGCTGGGGC AGGATTTCC	13860
AGTCTACCAA TGAAAATTCT CTATCCGGAG TTAGATGTGA CCATTATTGA TTCACTCAAAT	13920
AAGCGCATCA ACTTCCTACA ACTCTTGGCT CAAGAACTGG ATTTGAACGG AGTCATTT	13980
TACCACGGAC GTGCCGAAGA TTTTGCCCCAA GACAAGAACT TCCGTGCTCA ATATGATTT	14040
GTAACAGCTC GTGCCGGTTC CCGTATGCAG GTCCTATCTG AATTGACTAT TCCCTACCTT	14100
AAGGTTGGTG GCAAACATTG AGCACTCAAG GCTAGCAATG CGCCTGAGGA ATTATTAGAA	14160
GCTAAGAATG CCCTCAATCT CTTTTTTAGT AAGGTCGAAG ACAATCTCAG TACGCCCTAC	14220
CGAATAGAGA TCCCGCGCTAT ATCACAGTGG TAGAAAAGAA AAAAGAAACA CCAAATAAT	14280
ATCCACGTAA GGCTGGTATG CCAAATAAAC GCCCCACTTTA AATTTTTAG TAAACAAATG	14340
TTTACAAAAT CAGCCTCGCT CTTTTATTTTC TAGGCTCGGG AAAAATGAT TTACAAAATC	14400
AGCCTCGCTC TTTTATTTCT AGGCTCGGGAA AAAAATGATT TACAAAATCA TTTTTCTG	14460

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CTATACTATC CTAAGCAAAG GTTTTAATG TCATCCCGTG AGGTGACGAA GACGCAGAAA	14520
TATTTAAAC TCTTAAATTA CTAATTTTA AAGAAGTCCT ACTCTGAGGG CCTATTGCTG	14580
TAAAAATAATG GGCTCTTTTG TGATGCCAA AAGTGAGGTT TATATGAAAC AAGAATCAAC	14640
TGTTGATTTG TTAC	14654

(2) INFORMATION FOR SEQ ID NO: 107:

- (i) SEQUENCE CHARACTERISTICS:
- (A) LENGTH: 6405 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 107:

AGAAAAATCT GCTTACAGA AAATAAAAAT AATAGGAGAA AATCTATGTC AGATTTGAAA	60
AAATACGAAG GTGTCAATTCC AGCCTTCTAC GCATGTTATG ATGATCAAGG AGAAGTAAAGC	120
CCAGAACGTA CGCGTGCCTT GGTCAATAC TTCATTGATA AAGGTGTTCA AGGTCTTAT	180
GTCATGGTT CTTCTGGTGA ATGTATCTAC CAAAGCGTTG AAGATCGCAA GTTGATTTG	240
GAAGAAGTCA TGGCGGTAGC AAAGGTAAAT TGACCATTAT TGCCCATGTT GCTTGCAATA	300
ATACTAAAGA TAGTATGGAA CTTGCTCGCC ATGCTGAAAG CTTGGGAGTA GATGCTATTG	360
CAACGATTCC ACCAATTAT TTCCGCTTGC CAGAATACTC AGTTGCCAAA TACTGGAACG	420
ATATCAGTTC TGCGACTCCA AACACAGACT ACGTGATTTA CAACATTCTC CAATTGGCAG	480
GGGTTGCTTT GACTCCAAGC CTTTACACAG AAATGTTGAA AAATCCTCGT GTTATCGGTG	540
TGAAGAACTC TTCTATGCCA GTTCAAGATA TCCAAACCTT TGTCAGCCTT GGTGGAGAAG	600
ACCATATCGT CTTAATGGT CCTGATGAGC AGTTCTTAGG AGGACGCCTC ATGGGGCTA	660
GGGCTGGTAT CGGTGGTACT TATGGTGCTA TGCCAGAACT CTTCTTGAA CTCAATCAGT	720
TGATTGCGGA TAAGGACCTA GAAACAGCGC GTGAATTGCA GTATGCTATC AACGCAATCA	780
TTGGTAAACT CACTCTGCT CATGGAAATA TGTACGGTGT CATCAAAGAA GTCTTGAAAA	840
TCAATGAAGG CTTGAATATT GGATCTGTTG GTTCACCATT GACACCAGTG ACTGAAGAAG	900
ATCGTCCAGT TGTAGAAGCG GCTGCTGCCCT TGATTCGTGA AACCAAGGAG CGCTTCCTCT	960
AATCTAAAAG GAGGTATTTA TGACATATTA CGTTGCAATT GATATCGGTG GAACCAACAT	1020
CAAGTATGGT TTGGTTGATC AAGAGGGGCA ACTTCTTGAA TCGCATGAA TGCCAACTGA	1080
GGCGCATAAG GGTGGACCTC ATATCTTACA AAAGACAAA GATATCGTAG CTAGTTATT	1140
AGAAAAAGGC CCAGTAGCAG GTGTTGCCAT ATCTTCTGCT GGGATGGTGG ATCCGGATAA	1200

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GGGTGAGATT TTCTATGCTG GGCGCAAAT CCCTAACTAC GCAGGCACCC AGTTCAAAA	1260
GGAAATCGAA GAAAGCTTTA CTATTCCTTG TGAGATTGAA AATGATGTCA ACTGTGCAGG	1320
TCTTGCTGAG GCAGTATCTG GTTCAGGCAA GGGAGCAAGT GTGACACTTT GCTTGACCAT	1380
TGGAACCGGT ATCGGTGGTT GCTTGATTAT GGATAGGAAA GTCTTCCATG GTTTTAGCAA	1440
TTCAGCCTGT GAAGTCGGGT ATATGCATAT GCAGGATGGA GCTTTCAAG ACTTGGCTTC	1500
TACAACAGCT TTAGTGAAT ATGTAGCTGA AGCCCATGGA GAAGATGTTG ATCAGTGGAA	1560
TGGCCGTAGA ATTTTCAAAG AAGCCACTGA AGGAAACAAA ATCTGCATGG AAGGTATTGA	1620
CCGTATGGTT GACTATCTAG GAAAAGGTCT GCCAAATATT TGCTACGTTG CCAATCCAGA	1680
AGTGGTTATT CTTGGTGGTG CTATCATGGG GCAAGAGGCT ATCCTCAAAC CTAAGATCCG	1740
TACACCCTTG AAAGAGGCTT TGGTACCAAG TTTAGCAGAA AAAACACGAT TAGAATTGC	1800
CCATCACCAA AATACAGCAG GGATGTTGGG TCCATATTAT CATTAAAGA CAAAACAATC	1860
CTAGTTGGC TCAGCCAAAC TAGGATTTC TTACACGTTT TTGCTACGA TAGCCGTTGA	1920
GTTTTTATT TTCCCACTAG CTATTAAGA TTTTTCCCTT GCTTTCGCCA TTGATTTCCA	1980
AAAAGTAGGC ATAATCAAA TCGATAAAGA AGAGCATAGG AAGTTGAGCG GATATTGTT	2040
GGATATAGGA GGGTGGCTG TGGGTGGCTA CAAGAACAGT CTCTGTATAG GTCTGGCTAT	2100
CTTTATGGG AACACTGTA AAGAGTACAG TCTTGCCTT CATCTCCTTA GCATCTAATA	2160
GACTATCTAA AATAGAAGGA GTTGGCTG AAAGTGAGAA GCCCAGTACT AGACAATTTC	2220
CATCCATGAT GCTGGTTGTC CAGGCAAAGC CGTCTTGGTC TGTCAAAGCT TCGCAGACCA	2280
CACCTAGTCG CATAAAAGCT AATTCATTT CACGGGCGAC GAGGCCAGAA CTCCCTGTT	2340
CAAAGAGTA GATACGCTCA GCATCTCGA TTAGCTGGC AATTCGTTCT AGTTGGATT	2400
CGTCAATCAA GTCTGTGTT TGTCCCTCA TATTGCTATA ACTTCTGAGG ACTCGTTGG	2460
TCAGTGGACT GTGCTTGGAG ACTTGGTTGG CTTGATTTTC TGCCTGATGT TGGTATTGGA	2520
AAATAATTC TCGGTAGCCA GTAAAGCCAC ACTTTTCTGC AAAGCGGGTC AAAGCAGCTT	2580
GAGAAATATG TAATTTTGG GTGACTTCTT GAGAAGATAA ATCATCTGTA ATCCTTTCTAG	2640
CTTGCAAAAA ATAGCGAGCG ATTCTCTGTT CTAGGTCTGT CATTCTTC AATGTGAAT	2700
CAATGATAGT TCGGATATCT GGTTGTCCA TAGGGAAAGC TCCTTACAT GAGTCATACT	2760
CGAAGACTAG ATCAGAGAAT AGTCACACTT CATTATAACA CATAATATAA GGATAGATAA	2820
ATAAAAACGC ATCTCTGTT TAAAAACGAA AAAATCGAAA AAGCTTCTCT CTTTTCCATA	2880
ATTTCTACT CAAATTGTGG TACAATTAAG AGTAAGATTT TAAGTTAGAA ATGAGACTGA	2940

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TTTGTATGAG	AAAATTAAC	AGCCATTCGA	TTCCGATTG	GCTTAATTAA	TTGTTTCAA	3000
TCGTCATTT	ACTCTTTATG	ACCATTATTG	GTCGTTGTT	GTATATGCAG	GTTTTGAACA	3060
AGGATTTTA	CGAAAAAAAG	CTAGCTTCAG	CTAGTCAGAC	CAAGATTACA	AGCAGTTAG	3120
CCCCTGGGAA	AATTATGAT	GCTAGTGGAA	AACCTTTGGT	AGAAAATACG	TTAAAGCAGG	3180
TTGTTCCCT	TACCGTAGC	AATAAAATGA	CGGCTACAGA	CTTAAAAGAA	ACAGCTAAA	3240
AGTTACTGAC	TTATGTGAGC	ATCAGTTCTC	CAAATTGAC	AGAACGCCAG	CTGGCGGATT	3300
ACTATTTGGC	TGATCCTGAA	ATCTATAAAA	AAATAGTGGA	AGCTCTCCC	AGTGAGAAC	3360
GCTTGGATT	AGATGGCAAT	CGTCTATCCG	AATCAGAACT	GTATAACAAT	GCGGTCGATA	3420
GTGTACAAAC	GAGTCAACTA	AACTATACAG	AGGATGAAA	GAAAGAAATC	TATCTTTTA	3480
GTCAGTTAA	TGCTGTTGGA	AACTTTGCGA	CAGGAACCAT	TGCGACAGAT	CCTCTAAATG	3540
ATTCTCAGGT	GGCTGTTATT	GCCTCTATT	CAAAGGAGAT	GCCTGGCATT	AGTATTCTA	3600
CTTCTGGGA	TAGAAAGGTT	TTGGAAACCT	CCCTTCTTC	TATAGTTGGG	AGTGTATCCA	3660
GTGAAAAGC	TGGCTCCCA	GCGGAAGAAG	CAGAAGCCTA	TCTTAAAAAA	GGCTATTCTC	3720
TAAATGACCG	TGTAGGAACC	TCCTATTG	AAAAGCAATA	TGAAGAGACC	TTACAAGGAA	3780
AACGCTCGGT	AAAAGAAATC	CATCTGGATA	AATATGGCAA	TATGGAAAGC	GTGGATACAA	3840
TTGAGGAAGG	TAGTAAGGGA	AACAATATCA	AACTGACCAT	TGATTTGGCT	TTCCAAGATA	3900
GCGTGGATGC	TTTACTGAAA	AGTTATTCA	ATTCTGAGCT	AGAAAATGGT	GGAGCCAACT	3960
ATTCTGAAGG	TGTCATGCA	GTCGCCCTTA	ACCCAAAAAC	AGGTGCGGTT	TTGTCTATGT	4020
CAGGGATTAA	ACATGACTTG	AAAACGGGAG	AGTTGACGCC	TGATTCCTTG	GGAACGGTAA	4080
CCAATGTCTT	TGTTCCAGGT	TCGGTTGTCA	AGGCAGCGAC	CATCAGCTCA	GGTTGGGAAA	4140
ATGGAGTCCT	GTCAGGAAAC	CAGACCTTGA	CAGACCAGTC	CATTGTCTTC	CAAGGTTAG	4200
CTCCCCATCAA	TTCTTGGTAT	ACTCAGGCTT	ACGGTTCAATT	CCCTATCACA	GCGGTCCAAG	4260
CTCTGGAGTA	TTCATCAAAT	ACCTATATGG	TCCAAACAGC	CTTAGGTCTT	ATGGGGCAAA	4320
CCTATCAACC	CAATATGTTT	GTCGGCACCA	GCAATCTAGA	GTCTGCTATG	GAGAAACTGC	4380
GTTCAACCTT	TGGCGAATAT	GGCTTGGTA	CTGCGACAGG	AATTGACCTA	CCAGATGAAT	4440
CTACTGGATT	TGTTCCAAA	GAGTATAGCT	TTGCTAATT	CATTACTAAT	GCCTTGGGC	4500
AGTTTGATAA	CTATACGCCG	ATGCAGTTGG	CTCAGTATGT	AGCAACTATT	GCAAATAATG	4560
GTGTTCGTGT	GGCTCCTCGT	ATMGTGAAAG	GCATTTATGG	TAATAATGAT	AAGGGAGGAC	4620
TGGGTGACTT	GATTCAACAA	CTGCAACCGA	CAGAGATGAA	TAAGGTCAAT	ATATCCGACT	4680
CCGATATGAG	CATCTTGAC	CAAGGTTTT	ATCAGGTTGC	CCATGGTACT	AGTGGATTGA	4740

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CAACTGGACG TGCCTTTCA AATGGTGCCT TGGTATCCAT TAGCGGAAAA ACAGGTACAG	4800
CCGAAAGCTA TGTGGCAGAT GGTCAAGCAAG CAACCAATAC CAATGCGGTG GCCTATGCC	4860
CATCTGATAA TCCCCAAATC GCTGTCGCAAG TGGCTTTCC TCATAATACC AATCTAACAA	4920
ATGGTGTAGG ACCTTCCATT GCGCGTGACA TTATCAATCT GTATCAAAAA TACCATCCAA	4980
TGAATTAGAA AGGAAATTAT GCTTTATCCA ACACCTATTG CCAAGTTGAT TGACAGTTAT	5040
TCTAAAGTTAC CAGGTATCGG GATTAAGACG GCTACGCGTC TGGCCTTTA TACGATTGGG	5100
ATGTCTGCTG ATGATGTCAA TGAATTGCA AAAATCTCC TTTCTGCTAA GAGAGAATTG	5160
ACATATTGTT CTATTTGTGG ACGTTTGACA GACGACGATC CTTGTTCTAT CTGTAATGAT	5220
CCGACTCGTG ACCAGACAAC AATTTTAGTT CTTGAGGATA GTAGAGATGT GGCAGCCATG	5280
GAAAATATCC AAGAATACCA TGGACTCTAT CATGTCCTTC ATGGCCTCAT TTCTCCTATG	5340
AATGGTATCA GTCCGGACG TATCAATCTC AAGAGCCTTA TGACTCGTCT TATGGATAGT	5400
GAGGTTTCAG AAGTGATTGT GGCGACTAAT GCTACAGCGG ATGGTGAAGC GACTTCCATG	5460
TATCTTCAC GTTTGCTCAA GCCGGCTGGT ATCAAGGTTA CGCGTCTAGC ACGAGGTCTC	5520
GCTGTGGAG CGGACATTGA GTATGCGAC GAA GTGACAC TCTTACGAGC CATTGAAAAT	5580
CGGACAGAGT TGTAA GTGTA GGCAAATTTA CGAACTCCAT TCATTTATAA AAAATCAAAG	5640
AGGCTGAAAA TCGTTCTAT CGGGCTCTTT TTGTATAGTG TGATGAGTAG GCTCAGGTTC	5700
AACTTTAAA AAACCAAGCA AATATGATAT ACTAAAGAGC GAGTATTCTA GTAGAATTAG	5760
GACAAATAAT ATGAAACAAA CGATTATTCT TTTATATGGT GGACGGAGTG CGGAACGCCA	5820
AGTCTCTGTC CTTTCAGCTG AGAGTGTAT GCGTGCCTGC GATTACGACC GTTTCACAGT	5880
CAAGACTTTC TTTATCAGTC AGTCAGGTGA CTTTATCAA ACACAGGAAT TTAGTCATGC	5940
TCCGGGGCAA GAAGACCGTC TCATGACCAA TGAACCATT GATTGGATA AGAAAGTTGC	6000
ACCAAGTGCT ATCTACGAAG AAGGTGCAGT GGTCTTTCCA GTCTTCACG GGCAATGGG	6060
AGAAGATGGC TCTGTTCAAG GATTCTTGGA AGTTTGAAA ATGCCCTACG TTGGTTGCAA	6120
CATTTTGTC TCAAGTCTTG CCATGGATAA AACACAGACT AAGCGTGTTC TGGAAATCTGC	6180
TGGTATTGCC CAAGTTCTT ATGTGGCTAT CGTTGAAGGC GATGATGTGA CTGCTAAAAAT	6240
CGCTGAAGTG GAAGAAAAAT TGGCTTATCC AGTCTTCACT AAGCCGTCAA ACATGGGGTC	6300
TAGTGTGCGGT ATTCTAAGT CTGAAAACCA AGAAGAACTC CGTCAAGCCT TAAAACCTGC	6360
CTTCCGATAT GACAGCCGTG TCTTGGTTGA GCAAGGAGTG AATGC	6405

(2) INFORMATION FOR SEQ ID NO: 108:

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- (i) SEQUENCE CHARACTERISTICS:
- (A) LENGTH: 11309 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 108:

CGAGCTCGGG TACCGGATT TTAAGGAGTT TGATATGTAT AACCTATTAT TAACCATT	60
ATTAGTATTA TCTGTTGTGA TTGTGATTGC AATTTTCATG CAACCAACCA AAAACCAATC	120
CAGCAATGTA TTTGATGCCA GTTCAGGTGA TTTGTTTGAA CGCAGTAAAG CTCGCGTTT	180
TGAAGCTGTA ATGCAGCGTT TGACAGGGAT TTTAGTCTTT TTCTGGCTAG CCATTGCCTT	240
AGCATTGACG GTATTATCAA GTAGATAAGA AAATAATGGG CAGGACTAGG TCTTTGCC	300
TTTTTATTTT TAAAGGATGT TTGAGAAGGT TTTACAGTAA AAGAAAATTA AAAAATCTAG	360
AAAGAAAATA TGAAAGATAG AATAAAAGAA TATTTACAAG ACAAGGGAAA CGTGACTGTT	420
AATGATTGG CTCAGGCTT GGGAAAAGAC AGTTCCAAGG ATTTCGTGA GTTGATTAAA	480
ACCTTGCCT TAATGGAAAG AAAGCACCAA ATTCTGTTTG AAGAAGATGG TAGTCTGACA	540
TTAGAAAATTA AGAAAAAACA TGAGATTACC CTCAAGGGGA TTTTTCATGC CCATAAAAT	600
GGCTTGGCT TTGTTAGTCT GGAAGGCGAG GAGGACGACC TTTTTGTAGG GAAAATGAT	660
GTCAACTATG CTATTGATGG TGATACCGTC GAGGTAGTGA TTAAGAAAGT CGCTGACCGC	720
AATAAGGGAA CAGCAGCAGA AGCCAAAATT ATTGATATCC TAGAACACAG TTTGACAACA	780
GTTGTCGGGC AAATCGTTCT GGATCAGGAA AAACCTAAGT ATGCTGGCTA TATTGTTCA	840
AAAATCAGA AAATCAGTCA ACCGATTAT GTTAAGAAC CAGCCCTAAA ATTAGAAGGA	900
ACAGAAGTTC TCAAAGTCTT TATCGATAAA TACCAAGCA AGAAACATGA TTTCTTTGTC	960
GCGAGTGTTC TCGATGTAGT GGGACACTCA ACGGATGTCG GAATTGATGT TCTTGAGGTC	1020
TTGGAATCAA TGGACATTGT ATCCGAGTTT CCAGAAAGCTG TTGTTAAGGA AGCAGAAAAGT	1080
GTGCCTGATG CTCCGTCTCA AAAGGATATG GAAGGTCGTC TGGATCTAAG AGATGAAATT	1140
ACCTTACCA TTGACGGTGC GGATGCCAAG GACTTGGACG ATGCAGTGCA TATCAAGGCT	1200
CTGAAAATG GCAATCTGGA GTTTGGGTT CACATCGCAG ATGTTTCTTA TTATGTGACC	1260
GAGGGTCTG CCCTTGACAA GGAAGCCCTT AACCGTGCAG CTTCTGTTA CGTGACAGAC	1320
CGAGTGGTGC CAATGCTTCC AGAACGACTA TCAAATGGCA TCTGCTCTCT CAATCCCCAA	1380
GTTGACCGCC TGACCCAGTC TGCTATTATG GAGATTGATA AACATGGTCG TGTGGTCAAC	1440
TATACCATTA CACAAACAGT TATCAAGACC AGTTTCGTGA TGACCTATAG CGATGTCAAT	1500

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GATATCCTAG CTGGCGATGA AGAAAAGAGA AAAGAAATATC ATAAAATTGT ATCAAGTATC	1560
GAACTCATGG CCAAGCTTCA TGAAACTTTA GAAAACATGC CTGTGAAACG TGGAGCTCTC	1620
AATTTTGATA CCAATGAAGC GAAGATTTA GTGGATAAAC AAGGTAAGCC TGTTGATATC	1680
GTTCTTCGGC AGCGTGGTAT TGCCGAGCGG ATGATTGAGT CTTTTATGTT GATGGCTAAT	1740
GAAACAGTTG CCGAACATTT CAGCAAGTTG GATTTGCCTT TTATCTATCG AATTCAACGAG	1800
GAGCCTAAGG CTGAAAAGGT TCAGAAGTTT ATTGATTATG CTTCGAGTTT TGGCTTGCGC	1860
ATTTATGGAA CTGCCAGTGA GATTAGTCAG GAGGCACTTC AAGACATCAT GCCTGCTGTT	1920
GAGGGAGAAC CTTATGCAGA TGTATTGTCC ATGATGCTTC TTGCTCTAT GCAGCAGGCT	1980
CGTTATTCCGG AGCACAATCA CGGCCACTAT GGACTAGCTG CTGACTATTAA TACTCACTTT	2040
ACCAGTCAA TTCGTCGTTA TCCAGACCTT CTTGTTCACCA GTATGATTGCG GGATTACGGC	2100
CGTTCTAAGG AAATACCGAGA GCATTTTGAA CAAGTGATTC CAGAGATTGC GACCCAGTCT	2160
TCCAACCGTG AACGTCGTGC CATAGAAGCT GAGCGTGAAG TCGAAGCCAT GAAAAGGCT	2220
GAGTATATGG AAGAATACGT GGGTGAAGAG TATGATGCCAG TTGTATCAAG TATTGTCAAA	2280
TTCGGTCTCT TTGTCGAATT GCCAAACACA GTTGAAGGCT TGATTCACAT CACTAATCTG	2340
CCTGAATTTT ATCATTTCAA TGAGCGTGAT TTGACTCTTC GTGGAGAAAA ATCAGGTATC	2400
ACTTTCCGAG TGGGTAGCA GATCCGTATC CGTGTGAAA GAGCGGATAA AATGACTGG	2460
GAGATTGATT TTTCATTCTCGT ACCTAGTGAG TTGATGTTGA TTGAAAAGG CTTGAAACAG	2520
TCTAGTCGTA GTGGCAGAGG CGGTGATTCA AATCGTCGTT CGGATAAGAA GGAAGACAAG	2580
AGAAAATCAG GACGCTAAA TGATAAGCGT AAGCATTCAAC AAAAGACAA GAAGAAAAAA	2640
GGAAAGAAC CTTTTACAA GGAAGTAGCT AAGAAAGGAG CCAAGCATGG CAAAGGGCGA	2700
GGGAAAGGTC GTCGCACAAA ATAAAAAGGC ACGCCACGAC TATACAATCG TAGATACCGCT	2760
AGAGGCAGGG ATGGTCCTGA CTGGAACTGA AATCAAGAGT GTACGAGCTG CTCGAATTAA	2820
TCTCAAGGAT GGCTTGTCTC AAGTGAAAAA TGGAGAAGTT TGGCTGAGCA ATGTTCATAT	2880
CGCGCCTTAC GAAGAGGGCA ATATCTGAA CCAGGAACCA GAACGTCGTC GTAAACTCCT	2940
GCTCCATAAA AAGCAAATTC AAAATTTGGA ACAAGAGATC AAAGGGACAG GAATGACCTT	3000
AGTTCCCCTT AAGGTCTATA TAAAAGATGG CTACGCTAAG CTTCTTTAG GACTTGCCAA	3060
AGGGAAAGCAT GACTATGACA AACGGGAGTC TATCAAACGT CGTGAGCAAA ATCGAGATAT	3120
CGCGCGTGTG ATGAAAGCTG TTAATCAGCG ATAAAAAGAG GAATTGAAAAA TGGAAAAATT	3180
AGTTGCCTAT AAACGCATGC CTTTGTGGAA TAAACAAACAA ATGCCTGAAG CTGTTCAAGCA	3240

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AAAGCACAAAT	ACAAAAGTTG	GGACTTGCGG	GAAAATTACT	GTCTTGAAGG	GACCTCTCAA	3300		
GT	TATTGAA	TTGACAGAAG	AAGGGGAAGT	TCTAGCTGAA	CACCTCTTG	AAGCAGGGC	3360	
AGACA	ATGCCAAC	CTCAAGCCTG	GCACCGAGTG	GAAGCTGCCA	CAGATGATGT	3420		
GGAATGGTAC	TTGGAATT	ATTGTAAACC	TGAGGATTAT	TTTGCTAAAA	AATACAATAC	3480		
CAATCCTGTT	CATTCA	GAGG	TCCTAGAGGC	CATGCAGACA	GTGAAACAAG	GGAAAGCTT	3540	
GGATTTGGGT	TGTGGTCAGG	GGCGTAATT	TCTTTTCTA	GCCCAGCAAG	ATTTTGATGT	3600		
GACGGCTGTA	GATCAAAATG	GACTAGCTCT	TGAAATCTTG	CAAAGCATTG	TGGAGCAGGA	3660		
AGATTTGGAC	ATGCC	TGTTG	GCCTTACGA	TATCAATTCA	GCTAGCATTG	AAACAAGAATA	3720	
TGATTTTATC	GTTCAACAG	TTGTTCTCAT	GT	TTCTACAA	GCGGACCGCA	TTCCAGCTAT	3780	
TATTCAAAAT	ATGCAGGAGA	AAACCA	GTGTGTTAC	AACCTTATCG	TTTGTGCCAT	3840		
GGACACGGAG	GATTATCCTT	GCTCGTTAA	CTTCCCATT	ACCTTTAAAG	AAGGAGAACT	3900		
GGCAGACTAT	TACAAGGATT	GGGAATTGGT	TAAGTACAAT	GAAAATCCAG	GCCATTGCA	3960		
CCGTCGCGAT	GAGAATGGCA	ATCGTATTCA	ACTACGCTT	GCGACCTTAC	TAGCTAAGAA	4020		
AATCAAGTAA	ACACACATGA	AGATTAGGAA	TTTCC	CTGAT	CTTTTTCTT	TTTTACGAAT	4080	
GATATAGAAA	AGGAGGGAA	TCATGTTGT	TGCGAGAGAT	GCTAGGGAG	AATTGGTAAA	4140		
TGTGTTAGAG	GATAAACTTG	AGAGCAAGC	ATACACCTGC	CCAGCTTG	GAGGCCAGCT	4200		
CCATTTGCGT	CAAGGACCAA	GTGTACGGAC	GCAT	TTTGCC	CATAAATCCT	TAAAAGACTG	4260	
TGATTTTTTC	TTTGAAAATG	AAAGTCCAGA	ACACCTGGCC	AATAAGGAAT	CCCTCTATCA	4320		
CTGGTTGAAA	AAAGAGACAA	AGGTTCAATT	AGAGTACCCG	CTTCAGAAC	TTAAACAGAT	4380		
TGCGGATGTA	TTTGAAATG	GCAATCTAGC	TCTAGAAGTT	CAGTGTAGTC	CCTTGCCCTCA	4440		
GAAAGTCC	AAAGAGCGAA	GTGAGGGCTA	TCGTACTCAG	GGTTACCAAG	TACTGTGGTT	4500		
GCTGGTCAA	AAACTGTGGC	TCAAGGAGCG	TTTGACTCGT	CTACAGCAAG	GT	TTCTTTA	4560	
TTTCAGTC	AAACATGGGCT	TTTATGTTG	GGAATTAGAC	AAGGAAAAC	AAGTTTTAAG	4620		
ACTCAAATAC	CTGATT	TACC	AGGATCTCCG	CGGTAAAC	CATTATCAA	TCAAGGAATT	4680	
TTCCTATGGT	CAAGGTAGTT	TATTGAAAT	ATTGCGTCTT	CCCTATAAGA	GACAAAAAAT	4740		
ATCTCATT	TTT	ACAGTTCTG	AGGACAAGGA	CATCTCTCGC	TATACCGGC	AACAAC	4800	
TTATCAA	AAAT	CTCTTTGGA	TGAAAGAAC	AGCAGAAGCC	TATCAAAGG	GAGAAAATAT	4860	
CCTGACTTAT	GGACTGAAAG	AATGGTATCC	ACAAATT	CGA	CCAATAGTGG	GC	AAATTTT	4920
CCAGATTGAA	CAAGACTTGA	CTAGCTATT	TCAGCACTT	TATACCTATT	ACCA	AAAAAA	4980	
TCCTCAA	AAAT	GATTGGCAA	AGCTT	TATCC	ACCAGCCTT	TATCAGCAAT	ATTCTTGAA	5040

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AAATATGGTA	GAATAGAAG	GATGGAGGAA	TCTAATGGTA	TTACAAAGAA	ATGAAATAAA	5100
TGAAAAAGAT	ACATGGGATC	TATCAACGAT	CTACCCAACT	GACCAGGCTT	GGGAAGAAGC	5160
CTTAAAAGAT	TTAACAGAAC	AATTGGAGAC	AGTAGCCAG	TATGAAGGCC	ATCTCTTGGA	5220
TAGTGCAGAT	AACCTACTAG	AAATCACTGA	ATTTCTCTT	GAAATGGAAC	GCCAGATAGA	5280
GAAGCTTAC	GCTTATGCTC	ATATGAAGAA	TGACCAGGAT	ACACGTGAAG	CTAAGTATCA	5340
AGAGTACTAT	GCCAAGGCCA	TGACACTCTA	CAGCCAGTTA	GACCAAGCCT	TTTCATTCTA	5400
TGAGCCTGAA	TTTATGGAGA	TTAGCGAAAA	CGAGTATGCT	GACTTTTAG	AAGCTCAACC	5460
AAAGCTGCAG	GTTTATCAAC	ACTATTTGA	CAAGCTTTG	CAAGGCAAGG	ATCACGTTCT	5520
TTCACAAACGT	GAAGAAGAAT	TATTGGCTGG	AGCTGGAGAA	ATCTTGGTT	CAGCAAGTGA	5580
AACCTTCGCT	ATCTGGACA	ATGCGGATAT	TGTGTTCCCT	TATGTCCTAG	ACGATGATGG	5640
TAAAGAAGTT	CAGCTATCTC	ATGGGACTTA	CACACGTTG	ATGGAGTCTA	AAAAACGTGA	5700
GGTTGCCCGT	GGTGCCATTC	AAGCTCTTA	TGCGACTTAC	GAACAATTCC	AACACACCTA	5760
TGCCAAAACC	TTGCAAACCA	ATGTTAAGGT	GCAGGCTAAAG	TTCGTAACTA	5820	
CAAGAGTGC	CGTCATGCAG	CCCTCGCAGC	GAATTTGTT	CCAGAAAGTG	TTTATGACAA	5880
TTTGGTAGCA	GCAGTTCGCA	AGCATTGCC	ACTCTTACAT	CGCTATCTT	AGCTTCGTC	5940
AAAAATCTTG	GGGATTCAG	ATCTCAAGAT	GTACGATGTC	TACACACCGC	TTTCATCTGT	6000
TGAATACAGT	TTTACCTACC	AAGAAGCCTT	GAAAAAAAGCA	GAAGATGCTT	TGGCAGTCTT	6060
GGGTGAGGAT	TACTTGAGCC	GTGTTAACG	TGCCTTCAGC	GAGCGTTGGA	TTGATGTTA	6120
CGAAAATCAA	GGCAAGCGTT	CAGGTGCCTA	CTCTGGTGGT	TCTTATGATA	CCAATGCCTT	6180
TATGCTTCTC	AACTGGCAAG	ACAATCTGGA	CAATCTCTT	ACTCTTGTTC	ATGAAACAGG	6240
TCACAGTATG	CATTCAAGCT	ATACTCGTGA	AACTCAGCCT	TATGTTTACG	GGGATTACTC	6300
TATCTTTTG	GCTGAGATTG	CCTCAACTAC	CAATGAAAAT	ATCTTGACGG	AGAAATTATT	6360
GGAAGAAGTG	GAAGACGACG	CAACACGCTT	TGCTATTCTC	AATAACTTCC	TAGATGGTT	6420
CCGTGGAACA	CTTTCCGCC	AAACTCAATT	TGCTGAGTTT	GAACACGCCA	TTCACCAAGC	6480
AGATCAAAT	GGGGAGGTCT	TGACAAGCGA	TTTCCTAAAT	AAACTCTACG	CAGACTGAA	6540
CCAAGAGTAT	TATGGTTGA	GTAAGGAAGA	CAATCCTGAA	ATCCAATACG	ACTGGGCTCG	6600
CATTCCACAC	TTCTACTATA	ACTACTATGT	ATATCAATAT	TCAACTGGCT	TTGCGGCCGC	6660
CTCAGCCTG	GCTGAAAAAA	TTGTCCATGG	TAGTCAAGAA	GACCGTGACC	GCTATATCGA	6720
CTACCTCAAG	GCAGGTAAGT	CGGACTATCC	ACTTAATGTC	ATGAGAAAAG	CTGGTGTGGA	6780

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TATGGAGAAG	GAAGACTACC	TCAACGATGC	CTTTGCAGTC	TTTGAACGCC	GTTAAATGA	6840
GTTGAAGCC	CTTGTGAAA	AATTAGGATT	GGCATAAAAT	GGTTGAATCG	TATAGTAAGA	6900
ATGCTAACCA	TAACATGCGT	CGTCCTGTCG	TCAAAGAAGA	AATTGTAGAC	TTGATGCGTC	6960
AGCGTCAAAA	GCAGGTCACA	GGTTCTTGA	AAGAATTGGA	AGACTTGCC	CGCAAGGAAA	7020
ATATTCCAT	TATTCCCCAT	GAAACGGTTG	CTTATTTCCG	TTTCTTATG	GAAACCATGC	7080
AGCCTAAAAA	TATTCTGGAA	ATTGGGACGG	CTATCGGTTT	TTCAGCTCTC	TTGATGGCTG	7140
AACATGCGCC	AAATGCTAAG	ATTACAACTA	TTGATCGTAA	TCCAGAAATG	ATTGGTTTTG	7200
CCAAGGAAAA	TTTGCCCCAG	TTTGACAGTC	GCAAGCAAAT	CACTCTCCTA	GAGGGAGATG	7260
CGGTGGATGT	CTTATCTACA	CTGACAGAGT	CTTATGATT	CGTCTTTATG	GATTCTGCCA	7320
AGTCTAAATA	CATCGTCTTT	CTGCCAGAAA	TCCTCAAACA	TTTGAAGTT	GGTGGTGTGG	7380
TTGCTTGGG	TGATATTTTT	CAAGGTGGTG	ATGTTGCCA	GGATATTATG	GAAGTCGTC	7440
GTGGTCAGCG	AACCATTAT	CGAGGCCTTC	AAAAATTATT	TGATGCAACC	TTAGACAATC	7500
CAGAACTCAC	CGCAACATTA	GTGCCTTTAG	GAGATGGTAT	TCTCATGCTT	CGTAAAAATG	7560
TAGCAGATGT	TCAACTGTCT	CAAAGCGAAT	GATTTTCAGA	AAAATTTAAG	AAAAATAGT	7620
AAAATAGATA	GAGTAACACT	TATCTCAAAG	GAGTAGACAT	GAAGAAAAAA	TTATTGGCAG	7680
GTGCCATCAC	ACTATTATCA	GTAGCAACTT	TAGCAGCTTG	TTCGAAAGGG	TCAGAAGGTG	7740
CAGACCTTAT	CAGCATGAAA	GGGGATGTCA	TTACAGAAC	TCAATTAT	GAGCAAGTGA	7800
AAAGCAACCC	TTCAGCCAA	CAAGTCTTGT	AAATATGAC	CATCCAAAAA	GTTTTGAAA	7860
AAACAATATGG	CTCAGAGCTT	GATGATAAAG	AGGTTGATGA	TACTATTGCC	GAAGAAAAAA	7920
AAACAATATGG	CGAAAACCTAC	CAACGTGTCT	TGTCACAAGC	AGGTATGACT	CTTGAAACAC	7980
GTAAAGCTCA	AATTCTGACA	AGTAAATTAG	TTGAGTTGGC	AGTTAAGAAG	GTAGCAGAAG	8040
CTGAATTGAC	AGATGAAGCC	TATAAGAAG	CCTTTGATGA	GTACACTCCA	GATGTAACGG	8100
CTCAAATCAT	CCGTCTTAAT	AATGAAGATA	AGGCCAAAGA	AGTTCTCGAA	AAAGCCAAGG	8160
CAGAAGGTGC	TGATTTGCT	CAATTAGCCA	AAGATAATT	AACTGATGAA	AAAACAAAAG	8220
AAAATGGTGG	AGAAAATTACC	TTTGATTCTG	CTTCAACAGA	AGTACCTGAG	CAAGTCAAAAA	8280
AAGCCGCTTT	CGCTTTAGAT	GTGGATGGTG	TTTCTGATGT	GATTACAGCA	ACTGGCACAC	8340
AAGCCTACAG	TAGCCAATAT	TACATTGTAA	AACTCACTAA	AAAAACAGAA	AAATCATCTA	8400
ATATTGATGA	CTACAAAGAA	AAATTAAAAA	CTGTTATCTT	GAETCAAAAA	AAAATGATT	8460
CAACATTGT	TCAAAGCATT	ATCGGAAAAG	AATTGCAAGC	AGCCAATATC	AAGGTTAAGG	8520
ACCAAGCCTT	CCAAAATATC	TTTACCCAAT	ATATCGGTGG	TGGAGATTCA	AGCTCAAGCA	8580

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GTAGTACATC AAACGAATAG TCCAATCAA TGAGTCAGGG AAAAAACTCG ACTTCAGGAA	8640
AAAATGAAGC AAACATTCCC ACAATAAAC GCATAGTACA AGGTTTGTAC TGCCCCCAA	8700
AAAGTTAGAC AATTAATTAA TCCGAAGGAT TTAGTTCTGT ATTGCACAGA GCTAAGTCCT	8760
TTTAGTTTA TCTTAATTCT CTTATTGTTG TAATAATCAA TATAGTCTAT AATGGCTCGT	8820
TCCAATTGAT TAAGTGATTT AAATGTTTC TCATAGCCAT AAAACATTTG GGATTTAAA	8880
ATGCCAAAGA AAGATTCCAT CCTACCGTTG TCTTGGCTGT TGCCCTTACG TGACATGGAT	8940
GCTTGAATTG CCTTACTCTC TAGGAAGCGA TGATAAGAAT CGTGTGATA TTGCCAGCCT	9000
TGGTCACTAT GGAGAATCGT ATTCTCGTAG TGCTTCTCTT TGAATGCCCT TTCCAACATT	9060
AACGATCAAT CAATTTAACG ATGTACCTAA GATTAGAATT GTTTATCCCA AATTTATTTG	9120
AAAGCTTCTC TAAGCTATAT CCTTGTTCCTA TAAGTTCTA GATCTGAAC TTATCATCAT	9180
AAGTTAATT CATAATAAAA ACACCCCCAA AGTTAGATTT TTTCTGTCTA ACTTTGGGG	9240
TGTAGTTCAT GTACACCTGA TATGATGCGT TTTATAATT TAAAGACTTT TTGACCAGCC	9300
TCATTTTTT AACTTGATAC TCAGTGAAAA GCAAAGATTA AACTAGGAAG CTAGCTGTAG	9360
GCTGCTAAA GAACAGCTTT GAGGTTGTAG ATAAAACCTG TGAGGTCACC AACATATATA	9420
ATGTGAAGCT GACGTGGTTT GAATAGATTT TAGAAGAGTA TGAGTCTGGA AGTTTTAATG	9480
GATAATGCAA GATTCCATAG AATGGGTAAG CTAGAGTTCT TATGTGAAGA GTTGGGCAT	9540
AAACTTTTAC CTTTCTCTC CTACTCATCT TAGTATAGAA AAGTGAATCT GAAATAGTAC	9600
ATAACTGCTT CTAAAACATT CTTATAAATT GATTTAAATT CTCAAATCAT ATTATTCACT	9660
TCTTATTTC A TTTCTGTCTA CAATCCTGTT GAGAAGACAC GTGTTCATAT CAAAAGGTA	9720
TTGGCAAGTT GCAATACCTT TTTACGAGGC TCTGTTGTCT TATTTTTGTT TCAACTGACT	9780
ATATCTCCTA TGGTTCTAGT TCAGAAGGCT AGGCTATAAT TATGATTGAT AAGAAGTATC	9840
ATTCCAAGTA TTGGGAGTGA ATGTTCAAA ATCATGGGTT TCTATAATGG TCAGGCTGGC	9900
ATTTGCTAGA CGGCCATCTT TACGAAGAAC TGGTTCTTAA TAGCCTAGGA GAGTACGAAG	9960
ACTGGCAGTA AGATTGGCCG CGTGTCCGAC AATTAGAATA CGTTCAAGCTG GACTATCTT	10020
TAATGATTTG ATAAATGGG TGGTCCGTTG AGTTGACTA TAGAGGGATT CGGCTCCGAA	10080
CATTGAGTG TCAAATTGAG CAAGATTGAG ACGAAAAGCC TGGATTTGTT GCGGGTAAAT	10140
AGCTTCCAAG GTTGCATTT TCAAACCTTC TAACCTCCCA AGTTGCCATT CACGGAGATT	10200
AGGAACGATT TCTAAAGAAC AGGGGGTATA GAGTTGACTT TGGATAATCT CAGCAGATTT	10260
GACCGCTCGA GGTAAATCAC TTGAATAAT CTGATCAAA GGAATTTCCCT TGAGATACTG	10320

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ACCAAGTCGT TTTAGGGTTT CAATGGATTG AGGAAGAAGA GGAGAACAC CACTAGCACC	10380
TTGAAAACGA CCTCTTGGT TCCAGAGGGT ACGACCGTGG CGGACAAAGT AGAGTTTCAT	10440
TACTTGATGT CCTCCAAAAT ATCTACAAAG TCTGCCTTA CAAAGCTAGC CAAAGTCTTGT	10500
GGCGCGACGA TAATGCTGTG TCCGACTTCG CCTGCAGAGA CAATCATTTG ATCCAAATCT	10560
AGAGCAATTT TATCGATAAA AATGGGATAA TTGTGTTCT GACGAATTCC GACAGGATTA	10620
TTGGCTCCAT GAATGTAACC AGTTGTTTT TCTAAGTCCT TTTGTGGAAT CATGCTCACT	10680
TTTTTATTGC CAGAAATTTT AGCTAGTTTC TTTTCAGACA AGTGCTGAGT GATAGGGACA	10740
ATTCCGATAA TCGGTCCGGT CTTGTCTCCC AAAAGCGCCA AGGTTTGAA AATCTGATCT	10800
CGTTCATAAC CTTGAGGAAG CTCTCCTTCT AGGGCATTGA TTTGAATCCC CTGATGAGGG	10860
ATAGCTGCTT TAGATAGGAT TTGTTCCACC AATGTTTTT TGATTTAAC TTTTTTGCC	10920
ATTATTTATA TTTATCCTCC ATTGACTCA TCCAAATACC AAGCCAGATT CCCAGCGCAA	10980
AGAAGAAGGC GATGATGACA TAACCGACAA GTGAAAGTCC TGTGTATTGG ATACTTTCAG	11040
CGTTCCCTGC ATTTGGAATT AAGATCAAAA GGGTACTTGA TAGGACGATA CCGATGATGA	11100
AATGATAGAC GAACTGTTA CGGAGTTCTT CTAGTTCTCC GTCCGTCCAA GCGTAGGCCA	11160
CTTCTCTTT CTTGCCTTA CCTTTGGACA TCTTGAAAG AGGTGGGAGG GCAATATAGA	11220
CATGACCTGC CTCGACTAGC GGACGCATGT AACGGTAGAA AAATGTCAAG AGCAAGGTCT	11280
GGATATGGGC ACCGTCGGTA TCCGCATCG	11309

(2) INFORMATION FOR SEQ ID NO: 109:

- (i) SEQUENCE CHARACTERISTICS:
- (A) LENGTH: 5548 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 109:

CCATAGTCTA ACAAGTCTTT GTAAAGGTTT ATCCCTGATT CATGTAAGA TTGTGTAAG	60
AATCAAAAAA AGCCACTTTT GAAAAATGGC TGCTCCTAAA AATAGCTTAA AAAATTATTA	120
GTCCTGTGCG AAAGATTGGT TAGGAAGAAA AATCGTGAAG CAACTGCCTC TGCCAAGCTG	180
ACTCGTCACC GTGACTTGGC CACCTAATAA TTGACTGAGT TCTTTGACAA TGGCAAGGCC	240
AAGACCAGTG CCACCAAGTTT GTCTGCTTCG ACCTTTATTA ACTCGGTAAA AACGTTCAAA	300
AATACGATCC TGCTCTAATT GACTAATACC AATCCCTGTA TCTGATACAG AAATCTTAAT	360
GCCTTCGTTG ACCTTTGGG TCTTGACCTC AATTTTCCC CCTTGTTCAAG TGTAACGGAT	420

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GGCATTGGAT AAAAGATTGA GTAAGATTTG GGAAAGTAAT TGACTATCTG ATACGAGGGT	480
GACATCATCT GGCACCTGCA CCTTTAGCTG TAAATCCTTC TTCTTGAGCT GAGGTTGCAA	540
GCTTGTAGTC AAATCCTGTA CAAATTCTGC CAAAGAAGG GTCGTCCATT GTATAGGCAT	600
TTGTTGAGCC TTAGATAAGG TAAGAAGATG CTCAACAATA TGCTCAAGAC GCAAAACCTTC	660
TTTGTAAATA ATGTCTAGAA AGTCATCCTT GAGCGCTTCT TCTTCAGCTG ACATCCCCTT	720
AATGGTTCA GCAAAGCCCT TAATCGAAGT AACTGGTGTCT CTCATTCTAT GGGAGGCATT	780
TGAGACAAAG GCTAAATTAA ACTTTTCATA AGTTCTAAC TGTGTTAAAT CATATAGCAA	840
GACGAGCAC A GCTTCCACAG ATTGGGTGGG GCTAAAACG GGAACGTGCTG TCACATTCTAA	900
AATCAAGTCA CCCTCATGAA ACCCACTTAC TTCTTGTTTT AACCTTGTTT TTGATCAAA	960
GGCTTGGTGA ACTAAATTCC GAATATCCAT CCGTTTGAGG TCATCAAGTG AACTTATGTC	1020
CCCGTCCACA TCGGGAAAT AATGAGGCAG AGAGCGACTG GATAATAACA TCTGACCTTG	1080
AGCGGAAACT AAAAACGTCC CCATGGTTAG GTCCGACAGA AGAACCTCCA TTGTTTCCGC	1140
TAGATCCTTG TATTGCTGAT CCTGTTGGG GACTTTGGTT TTTAGGCCAG ACACATACTG	1200
AGCCAAGAC TTTAAGTCTT CTTGCCCTT TTCTAAAAAG TATTCACTAC TGGTCAAGAG	1260
AGGTTGGTGC AAGGTCTCAA AAGCAACTTC CCATTTCAGA AGGCAAAAGA GCCAGTAGCC	1320
ACCTAGTCCC AAAGAAAGGG CTAGAAGAAA GAGACCGATG CCTTTACTGA TCCAAGTTAA	1380
TGCCATCCCT GCAATCAGAA TGAGGCTAAC ACTTAGATTG ACTAGCCAAA ATTGAAGGTA	1440
GGGTTCATC TATAACTCCT TGAACCTATA ACCATAACCC CGAATGGTTC GAATAAATTG	1500
AGGGGCTTTA GGATTGCTTT CAATTTCCTT CCTCACTTA CCAATATGAA CGTCCACCAA	1560
ACGTGTTCC TGCCCAAAGT CATACCCCCA GATACGTTCC AAAAGACGCT CTCTAGTCAG	1620
TGTCATGTTG GGATGTTCA TAAGATAGAG CAAGAGTTCA AATTCTTTG GGGTCAAAC	1680
CAGTAACCTA TTCCGCTTGT AGACTTCATG ACGCTCAGGG TATACTTCA AGGTCCAAA	1740
TAGCCAAGAA TCGTCAGCGA TATTATCTGA ATCATCTCCT TCTTGTTCTC CTTTAGTTCG	1800
CCTGAGGACA GCCTTGACAC GCGCCAGCAA TTCTCTAGGG CTAAAAGGCT TGGTCAGGTA	1860
GTCATCAGCC CCTAATTCCA AGGCCAAAAC CTTATCAAAT TCATCACTT TCGCAGAAC	1920
CATCATAATT GGAGTTTGA CCCCTTTGGC TCTCAGCCGC TTACAAACTT CCATGCCATC	1980
TAATTGTTGGT AACATGATAT CAAGCAAGAT AAAATCAAAG GGTCTGTTT CTGCCAAAGC	2040
TAAGGCCTTC CGTCCATTG TCACCAATTG AGTAGAAAAG CCTTCCTTAC TTAAATGGTA	2100
GTCAAGCAAT TTCAGAATGT GTTCTTCATC ATCCACTAAT AAGACTTGGTT TTGTCATCTA	2160

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TTATCCCTA TTGTAACAT TATAACACAA TTATCAGAAA TCCTAACATT GCTAAATCAG	2220
ATTAATTTG CCTATCAAGA CTAGTATCTG GTCAAACGCT CAATCATCTC CTTGTGCTCT	2280
GGATAGTCG CCAGTAGATC TACCTTTCA AATAATTCAA ATCCTCAA TTCAAAACCA	2340
GGAGCAACAA GACAAGAAC CAGAGCATCA TCCTTATCAA CTGTTGATCC CCAAATAGTG	2400
CCCTTAGGAA CACAGTAGTG AAGTTGTTGC CCTTTGGATA TGTCCAGGCC TAAAGTGA	2460
GCTTCGTAGT GACCATCTGC TGTAAATCATG TGAACAGTAA GTGGGGATCC TGCATGAAAA	2520
TACCAAGATTT CATCTGCTGT CAATCGGTGA AAATGTGAAG GATTGTTTC TTCTAATAAG	2580
AAATAAAATAC TGGTATAAAAG CGCCCTTCCC TTACCAAGCAA GGTTTATAGT GTCTGAAGCT	2640
TTTTTGTTT GTCTAAAATA GCCACCTTCA ATATGGGGAG CTAACCTCTAG AGTTCTTATC	2700
AAGTCTTCTT TATCCGTCGG AGCCAATGGG TTGAAGTAAC TCTTGTCAA AGTGGTTTA	2760
CGATTCAAG AACCCCTCTC AGTTCTGAGG ACACGGTAAT GATTGATGCC ACGGAAGTAC	2820
AAATCAATCG CCCTAAAAAA AGAATTAGCG AATGATTCTG GTAAAAAAAA TGCCACGCTA	2880
TGAAGGCTCA AGCGATTGTC ACAAGTCAAG GGAGAATTGT TTCTTTGGAT ATCGCTGTGA	2940
ACTATTGTCA TGATATGAAG TTGTTCAAAA TGAGTCGCAG AAATATCGGA CAAGCTGGTA	3000
AAATCTTGGC TGACAGTGGT TATCAAGGGC TCATGAAGAT ATATCCTCAA GCACAAACTC	3060
CACGTAATC CAGCAAACCTC AAGCCACTAA CAGTTGAAGA TAAAGCCTAT AACCATGCC	3120
TATCCAAGGA GAGAAGCAAG GTTGAGAACAA TCTTTGCCAA AGTAAAAACG TTTAAAATGA	3180
TTTCAACAAAC CTATCGAAAT CATCGAACAC ACTTCGGATT ACGAATGAAT TTGATTGCTG	3240
GCATTATCAA TCATGAACTA GGATTCTAGT TTTGCAGGAA GTCTATTATT TGGTTAGGTG	3300
AATTAGTGAAGC GCGTTAGGC AAGTGTCTCT GGTTACGACG TCATGGACTC TAAATCGATT	3360
ATATTTAGGG GTCATGACTA GTGAAGCAGT TAGCTAGTTC GCATATAAGC GGCTAGCGTC	3420
TAACAAATTAG GAACCTTACTG TCCAATAACT TTAAGATTAC GACGTTTAG GACATAAATC	3480
GATCATATTT ATGCTCTAAA ACTAGTGAAG CGCCTAGCCA AAGTCCGAAT AGGATTTGGC	3540
GTTAGTTACT TAGATTGCTT TGCAATCAAG TAACTTTGGC GATTTACATC TTCTCTGGCG	3600
CTTCTACTCC AAGCAAGCGA AGGGCTTCTT TGAGAACGAC TGCGGTTGCC TAGCTGAGGG	3660
CTAGACGGCT GTCGCGTTCT GGGCTTCCAT CCAAGATACG TGTATGTGCA TAGTATTGTT	3720
TAAAGGATTG AGCCAGGCTA ATTGCAAATT TAGCAATGAT AGAAGGTTCA AAGTTATCTG	3780
CCGCACGGTT GATAATACGT GGGAACTCTT GAATGAGTTT AATGATTTCC CAGCTTTAG	3840
TATCATTCAA GCTATAGTTG CCAGCTGTTT CTGGTTGAA ATCGGCTTG CGTAAGATAG	3900
ATTGGATACG AGCGTAGGCA TATTGAACGT AAGGTCCAGT TTCACCCCTCG AAGGATACCA	3960

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TAGCCTCTAG	GTCGAAGTCG	TATCCATTG	TACGGTCGGT	TTTGAGGTCA	TAGAATTAA	4020
TGGCTCCAAT	CCCAACAGCA	TGTGCTACTT	GGTCTTGT	TTCTAGTTCA	GGATTTTAG	4080
CCTCGATTTG	GACCTTGGCA	CGGCTAACAG	CCTCTGCAAC	AGTAGGCTCT	AGCAAGATGA	4140
CATTCCTTT	ACGAGTAGAG	AGTTTCTTCC	CTTCTTTGT	AACCAAACCA	AAAGGAACGT	4200
GAGTAATGTC	GTCACTCCAG	TCGTAGCCC	TCTCTTCAA	GACAGCTTG	AGCTGTTAA	4260
AGTGGGCAGA	TTGTTCTTGA	CCAAACGACAT	AGATAGATT	AGCAAATTGG	TATTCGTTTT	4320
TACGGTAGAG	GGCTGCAGCC	AAGTCACGTG	TGATATAGAG	AGTTGCACCA	TCAGACTTCT	4380
TGATGAGGGC	TGGATGTTCA	ATTCCATATT	TCTCAAGATT	CACAACCTGG	GCACCTTCTG	4440
ATTCAAGAAG	TAGTCCTTT	TCAGAAAGAA	TGTCTACAAC	TGCATCCATC	TTATCATTGT	4500
AGAAGGCTTC	TCCGTTATAG	CTGTCAAATT	CAACCTCAA	TTCATTGTAA	AGGCGGTTAA	4560
ATTCCACTAA	ACTTTCATCG	CGGAACCATT	GCCAAAGAGC	GAGAGCTTCC	TCATCTCCAT	4620
TTTCAAGTTT	ACGGAACCATT	TCGCGCGCTT	CTTCATCCAA	GCTAGGGTCA	TTTCAGCTT	4680
CAGCGTTGAT	GCGGACATAG	AGTTTAAGGA	GTTCATCGAT	TGGATGACCT	TTTACAGCTT	4740
CTTCGTCGCC	CCATTTTTG	TAGGCAACAA	TCAACATCCC	AAATTGTTA	CCCCAGTCTC	4800
CCAAATGGTT	GACCTTGACC	GTGATAAAC	CGATTTTTG	AAAAATATGT	GACAAGCTAT	4860
CTCCGATAAC	AGTTGAACGC	AGGTGGCAA	TAGAAAATGG	TTTAGCGATA	TTGGACTAG	4920
ACATGTCGAT	AACAAACATTT	TCTTGTAC	CAATTTTG	GTCAGCATAG	TGTTCTTTT	4980
CAGTGGTAAC	AGCTTGCAAT	ACTTGAGCAG	AAATGGCAGA	TTTATCAAGG	AAAAAGTTAA	5040
CGTAAGGTCC	TGTTGCGACA	ACTTTTCAA	AGGCTGGCT	GTTCATTTT	TCAGCCAGTT	5100
CAGCCGAAT	CATTGTTGGT	GCTTACGTT	CGACTTTGC	AAGAGAAAAA	GCAGGGAAAG	5160
CAATGTCCTC	CATTCTGAG	TTTTTAGGGG	TTTCCAGTAA	CTTTAAAATA	GCCTCTGGT	5220
CCAGGCTATC	AATGATGCTA	GATAATTGCG	TAGCAATCAA	TTCTTTGTA	TTCATTAAGA	5280
GCTCCTTTT	GGACTTTCT	ACTATTTAT	CACAATTAA	AAGAAAGAAG	AAAAAATTTT	5340
TGAAATCTCC	TGTTTTTTG	GTATAATATG	GTTATAAATA	TAGTTATAAA	TATGCACCGCA	5400
AGAGGATTTT	ATGAGAAAAA	GAGATCGTCA	TCAGTTAATA	AAAAAAATGA	TTACTGAGGA	5460
GAAATTAAAGT	ACACAAAAAG	AAATTCAAGA	TCGGTTGGAG	GCGCACAATG	TTTGTGTGAC	5520
GCAGACAACC	TTGTCCTCGTG	ATTTGCAGG				5548

(2) INFORMATION FOR SEQ ID NO: 110:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 3132 base pairs

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- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 110:

TACCCGGTAG TCTTAGCAGA CACATCTAGC TCTGAAGATG CTTTAAACAT CTCTGATAAA	60
GAAAAAGTAG CAGAAAATAA AGAGAACAT GAAAATATCC ATAGTGCTAT GGAAACTTCA	120
CAGGATTTA AAGAGAAGAA AACAGCAGTC ATTAAGGAAA AAGAAGTTGT TAGTAAAAT	180
CCTGTGATAG ACAATAACAC TAGCAATGAA GAAGCAAAAA TCAAAGAAGA AAATTCCAAT	240
AAATCCAAG GAGATTATAC GGACTCATTT GTGAATAAA ACACAGAAAA TCCCAAAAAA	300
GAAGATAAG TTGTCTATAT TGCTGAATTT AAAGATAAAAG AATCTGGAGA AAAAGCAATC	360
AAGGAACAT CCAGCTTAA GAATACAAAA GTTTATATA CTTATGATAG AATTTTAAC	420
GGTAGTGCCA TAGAAAACAAC TCCAGATAAC TTGGACAAAA TTAAACAAAT AGAAGGTATT	480
TCATCGGTTG AAAGGGCACA AAAAGTCCAA CCCATGATGA ATCATGCCAG AAAGGAATT	540
GGAGTTGAGG AAGGTATTGA TTACCTAAAG TCTATCAATG CTCCGTTGG GAAAATTTT	600
GATGGTAGAG GTATGGTCAT TTCAAATATC GATACTGGAA CAGATTATAG ACATAAGGCT	660
ATGAGAATCG ATGATGATGC CAAAGCCTCA ATGAGATTTA AAAAAGAAGA CTTAAAAGGC	720
ACTGATAAAA ATTATTGGTT GAGTGATAAA ATCCCTCATG CGTTCAATTA TTATAATGGT	780
GGCAAAATCA CTGTAGAAAA ATATGATGAT GGAAGGGATT ATTTTGACCC ACATGGGATG	840
CATATTGCAG GGATTCTTGC TGGAATGAT ACTGAACAAG ACATCAAAAA CTTTAACGGC	900
ATAGATGGAA TTGACACCTAA TGACCAAATT TTCTCTTACA AAATGTATTG TGACGCAGGA	960
TCTGGGTTTG CGGGTGTGAA AACAAATGTTT CATGCTATTG AAGATTCTAT CAAACACAAC	1020
GTTGATGTTG TTTCGGTATC ATCTGGTTT ACAGGAACAG GTCTTGTAGG TGAGAAATAT	1080
TGGCAAGCTA TTGGGCATT AAGAAAAGCA GGCATTCCAA TGGTTGTCGC TACGGGTAAC	1140
TATGCGACTT CTGCTTCAAG TTCTTCATGG GATTAGTAG CAAATAATCA TCTGAAAATG	1200
ACCGACACTG GAAATGTAAC ACCGAACGTCA GCACATGAAG ATGCGATAGC GGTCGCTTCT	1260
GCTAAAATC AAACAGTTGA GTTTGATAAA GTTAACATAG GTGGAGAAAG TTTTAAATAC	1320
AGAAAATATG GGGCCTTTT CGATAAGAGT AAAATCACAA CAAATGAAGA TGGAAACAAA	1380
GCTCCTAGTA AATTAAAATT TGTTATATATA GGCAAGGGGC AAGACCAAGA TTTGATAGGT	1440
TTGGATCTTA GGGCAAAAT TGCAGTAATG GATAGAATTT ATACAAAGGA TTTAAAAAT	1500
GCTTTAAAAA AAGCTATGGA TAAGGGTGCA CGCGCCATTA TGGTTGTAAA TACTGTAAAT	1560

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TACTACAATA GAGATAATTG GACAGAGCTT CCAGCTATGG GATATGAAGC GGATGAAGGT	1620
ACTAAAAGTC AAGTGTTC AATTCAGGA GATGATGGTG TAAAGCTATG GAACATGATT	1680
AATCCTGATA AAAAAGTGA AGTCAAAAGA AATAATAAAG AAGATTTAA AGATAAATTG	1740
GAGCAAACT ATCCAATTGA TATGGAAAGT TTTAATTCCA ACAAAACGAA TGTAGGTGAC	1800
GAAAAAGAGA TTGACTTTAA GTTGCACCT GACACAGACA AAGAACTCTA TAAAGAAGAT	1860
ATCATCGTTC CAGCAGGATC TACATCTTGG GGGCCAAGAA TAGATTTACT TTTAAAACCC	1920
GATGTTTCAG CACCTGGTAA AAATATTAAA TCCACGCTTA ATGTTATTAA TGGCAAATCA	1980
ACTTATGGCT ATATGTCAGG AACTAGTATG GCGACTCCAA TCGTGGCAGC TTCTACTGTT	2040
TTGATTAGAC CGAAATTAAA GGAAATGCTT GAAAGACCTG TATTGAAAAA TCTTAAGGGA	2100
GATGACAAAA TAGATCTTAC AAGTCTTACA AAAATTGCC CACAAATAC TGCGCGACCT	2160
ATGATGGATG CAACTCTTG GAAAGAAAAA AGTCAAACT ACT TTGCATCACC TAGACAACAG	2220
GGAGCAGGCC TAATTAATGT GGCAATGCTT TTGAGAAATG AAGTTGTAGC AACTTTCAA	2280
AACACTGATT CTAAAGGTTT GGTAAACTCA TATGGTTCCA TTTCTCTTAA AGAAATAAA	2340
GGTGATAAAA AATACTTAC AATCAAGCTT CACAATACAT CAAACAGACC TTTGACTTT	2400
AAAGTTTCAG CATCAGCGAT AACTACAGAT TCTCTAACTG ACAGATTAAA ACTTGATGAA	2460
ACATATAAG ATGAAAAATC TCCAGATGGT AAGCAAATTG TTCCAGAAAT TCACCCAGAA	2520
AAAGTCAAAG GAGCAAATAT CACATTTGAG CATGATACTT TCACTATAGG CGCAAATTCT	2580
AGCTTGATT TGAATGCGGT TATAATGTT GGAGAGGCCA AAAACAAAAA TAAATTGTA	2640
GAATCATTAA TTCATTTGA GTCACTGGAA GCGATGGAAG CTCTAAACTC CAGCGGGAAAG	2700
AAAATAAACT TCCAACCTTC TTTGTCGATG CCTCTAACTG GATTTGCTGG GAATTGGAAC	2760
CACGAACCAA TCCTTGATAA ATGGGCTTGG GAAGAAGGGT CAAGATCAA AACACTGGAA	2820
GGTTATGATG ATGATGGTAA ACCGAAAATT CCAGGAACCT TAAATAAGGG ATTGGTGGAA	2880
GAACATGGTA TAGATAAATT TAATCCAGCA GGAGTTATAC AAAATAGAAA AGATAAAAAT	2940
ACAACATCCC TGGATCAAAA TCCAGAATTAA TTGCTTCA ATAACGAAGG GATCAACGCT	3000
CCATCATCAA GTGGTTCTAA GATTGCTAAC ATTTATCCTT TAGATTCAA TGGAAATCCT	3060
CAAGATGCTC AACTGAAAG AGGATTAACA CCTTCTCCAC TTGTATTAAG AAGTGCAGAA	3120
GAAGGATTGA TT	3132

(2) INFORMATION FOR SEQ ID NO: 111:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 14672 base pairs

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- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 111:

CGAGATTTCT TTAAATGAAC TACGTGAAAT CTACCCATCA TCCAGATCTG GATATTCTCT	60
CCTATCTATA AGTAAAGTT TAGGAGATTT TAATATAAGT TCTCATGCTT TTAAAGCTTC	120
GGTAAGAGAT TTAAAACCGC TCAGTTCCC ACTCATTTGC TTCTGGGAGA GTTCTCATT	180
TATTATTCTT GAAAAAATTA GTAAAAACAA GTTTTATATT TTAGATCCTG CAAAAGGCAG	240
GCAGAGAATG TCAATAAGTG AATTGAAAG GCATTATTCA AATATCATT TAACATTTAA	300
AAAGTTAGAT AGCTTTATGT CTCGTAAAGA TAATAAGAAG TCGCCTGTT TAAAGTATT	360
TTTTAAGTAT AGGAATAAGC TAGGGATTTT ATTTTTGTA ACAGCATTAT TGTATGTAAT	420
ACAATCATT A GTACCTATAG CTAATAGATA CATAATTGC ACGAATTTCAG AGGACGATT	480
GTATTCGTCT AGAATGTTAT TTACTATATT ATTTATATT ACTGTTTCAT TCTCACTAAT	540
GTATTTATTA AGACAGATAT ATGTTGCATC CTTAAAATAT ATAATGGATA AAGAGATTAG	600
CTATGATTTT ATGAAACATT TGATATATT ACCTTACAGT TTTTATGAAA AACGTACTTT	660
AGGGGATATA CTTTTAGAG CTAACTCTAT TGTTTATATA AGAGAAATAC TATCAAATAA	720
TTTTATAGCA GCTATACTTG ATTTGTTAAT GATTGTGGTT TATGCTGTGG TTTTATTTAG	780
CTTTTCTAAG TACATGGTAA TCTTTTAAT ATCACTAAGT CTAGCTCTAT CTATTGTAAT	840
GTATCCAATC ATAAAAATCT CAAAAAAATT AATTGATAAA AATATAAAAG AAAAGGTTAA	900
TGTTCAAAAT ATTACTTCGG AAGTAATTTC TAAAAATAGT GATATTAAGC TAACTGGAGA	960
AGAGGAATT TGGATTAACA AATGGGATAA TTTAATACA AAACAGCTCA TCATAGGTGG	1020
AAAACCTGAT ATACATTAT CAATTGTTAG TAGTATAACG AATGTTTTAC AAATTATTCT	1080
CCCTGTTTG ACCCTTATTG TAGGTGTAA TATAAAACA TTCGAACAAT TGACGTTAGG	1140
ACAAATTGTA GCAATAAGTA CAGTCTCACC ATACTTTATT TCTCCTATAA TTTCTTTAAG	1200
TGATAACTAT ATACAATTAA TGTTATTAAA GGGATATT TTAAAGAATAG AGGATGTGTT	1260
TAATACTAAA TCCGAATTAA TTCCAGAAAG AGTCAGTCAA GATATAAAAT TTGATAAAAA	1320
AATAGAATTAA AAAGATATTG GGTATAAATA TGGATTATTG GATGATTATG TTTTGAAAGG	1380
AATAAAATGTT ACTATTTAAA AAGGAGAAC TGTGCTATT GTGGAGAAT CAGGTTCAAGG	1440
TAAGAGTACA TTAGCTAAA TTTTATTAGG TTTATTAGAA CCTAATATTG GTTCAATAGA	1500
AGTTGATGGA GTAGAAAAAG AAGAAATTGG TCAAACATTG TATAGAAAGA TTTTTGGAGC	1560

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AGTGTACAA AATTCAACCC TAAGTTATGG TACCTTAAGA GAGAATTGGA CATTGGACA	1620
CTTTGTTCA GATGAAGAAT TAATGACAAA TCTAAATTCA ATTGGTCTTA GCAATGTAGT	1680
TAAATCTTTA CCTCTTGAT TAGAGACAAT CATCGCTGAA GAAGGTAATA ACTTTCTGG	1740
AGGCCAGCAG CAAATGATAC TTTTAGCTCG TTGTCCTTG TCGAAACCTT CGGTAGTTGT	1800
TTTGGACGAA GCAACAAGTA GTT TAGATAA TTTATCTAA CAAATTACAA CTTCTTACTT	1860
AAGTGAATC GGTACCACTA AGATTTAAC TGCCCATCGA CTAGATACTA TCAAGTCTGC	1920
AGATAAGATC TTAGTAATGC ATAATGGTGA AATTGTAGAG ATTGGGACCC ATAGAGAACT	1980
TCTTGACTA GGAGGCATTT ATAAGCAATT CTATTCAAAT AATTAGTTTG TGATTTAAAG	2040
GGTAAATTAA TGAAGATTAT GAAAAAAA TATTGGACTT TAGCGATATT ATTCTTTGT	2100
TTGTTCAATA ATTCTGTTAC TGCTCAAGAA ATACCTAAA ATCTTGATGG CAATATAACT	2160
CACACTCAGA CTAGCGAAAG TTTTCTGAA TCTGATGAAA AACAGGTTGA CTATTCTAAT	2220
AAAAATCAAG AAGAAGTGA CCAAAATAAA TTTCGTATTC AAATCGATAA GACAGAATTA	2280
TTTGTAAACAA CAGATAACAA TTAGAAAAA AACTGTTGTA AATTGGAAC TGAACCACAA	2340
ATAAAATAACG ATATTGTTAA CTCTGAAAGT AATAATTAC TAGGCGAAGA TAATTTAGAT	2400
AATAAAATTA AGGAAAATGT TTCTCATCTA GATAATAGAG GAGGAAATAT AGAGCATGAC	2460
AAAGATAACT TAGAATCGTC GATTGTAAGA AAATATGAAT GGGATATAGA TAAAGTTACT	2520
GGTGGAGGCG AAAGTTATAA ATTATATTCT AAAAGTAATT CTAAAGTTTC AATTGCTATT	2580
TTAGATTCA GAGTCGATTT ACAAAATACT GGATTACTGA AAAATCTTC AAATCACTCA	2640
AAAAACTATG TCCCCAATAA AGGATATTAA GGAAAGAGG AGGGAGAGGA AGGAATAATA	2700
TCAGATATTCA AAGATAGATT AGGTCACTGG ACAGCTGTTG TAGCTCAAAT TGTAGGGAT	2760
GACAATATTA ATGGAGTAAA TCCTCACGTT AATATTAACG TCTATAGAAAT ATTTGGTAAG	2820
TCGTCAGCTA GTCCAGATTG GATTGTAAAA GCAATTGTTG ATGCTGTAGA TGATGGCAAT	2880
GATATTATCA ATCTTAGTAC TGGACAATAT TTAATGATTG ATGGAGAATA TGAGGACGGA	2940
ACAAATGATT TTGAAACATT TTGAAAGTAT AAAAGGCTA TTGATTACGC GAATCAAAA	3000
GGAGTAATTA TAGTAGCTGC ATTAGGAAAT GACTCCCTAA ATGTATCAA TCAGTCAGAT	3060
TTATTGAAAC TTATTAGTTC ACGCAAAAAA GTAAGAAAAC CAGGATTAGT AGTTGATGTT	3120
CCAAGTTATT TCTCATCTAC AATTTGGTC GGAGGCATAG ATCGCTTAGG TAATTTATCA	3180
GATTTTAGCA ATAAAGGGGA TTCTGATGCA ATATATGCGC CTGCAGGCTC AACATTATCT	3240
CTTTCAAGAT TAGGACTTAA TAACTTTATT AATGCGAGAA AATATAAAGA AGATTGGATT	3300

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TTTTCGGCAA CACTAGGAGG ATATACGTAT CTTTATGGAA ACTCATTGCG TGCTCCCTAAA	3360
GTTTCTGGTG CGATTGCAAT GATTATTGAT AAATACAAAT TAAAAGATCA GCCCTATAAT	3420
TATATGTTG TAAAAAAATT CTGGAAGAAA CATTACCACT AAAAAATGGT ATAAAAGTGT	3480
TAAATATACC AAACGTATTG AGATATGATT TGAATATGTT ACAATTAGAA TATAAAATG	3540
AACAAAGTTG GGATAGTTTC ATAGATAATG TTAATTAAAT TGAGTTGGAA GAGAGAATTC	3600
AAACTACTAT TGGAATTAAA CAAATAAACAA CACACAATAT TATTACTATT CCCCGAGAAG	3660
GGTACTCTCA AAATTATTTA CCTAACACTT CAGAAATAC ATATAATTCA TTACAAGTCA	3720
GTTTAGTTGG AGTATTACTA CTTTTATATAA GTATGGTAA TATTTATGG CCTAAAAAAA	3780
GTAAATGAAA ATAAAATTG GAGCCCTCTG AAAAGTAAG TCCTACAGTT CAACTAAAAT	3840
GAGTCAAAAG ATGAATCACC TTGATGTAGG GGAGTTGTGTC TTATTGCTGC CTGAACACCT	3900
CCGTTCAAGAG GAAGAACATT ATAAAATCTGT TTTTGAAGAC GACTTAACCA GTCGCATATC	3960
TAGTCAAGAT GAACGACAGC AAATGACTGC TACGGTAGGT TATTTAGAAT CAGGTCAGGA	4020
TCGTTTGTC TATAATACGA CCCCTATTC TTACCAAGCAG TTTTGAAAG ATCCAATCAT	4080
CATTGTTATA ACACCCCAAT CAACTGGTCC ACAGTCATT TTGTTTTGGAA TAGACGCAGT	4140
ACAGAACTAC GTTCTCTTTA ATCAATTGTC TGATGCCAG GAGCTTATCC AGAGACAAGG	4200
CATTGAAAAT TGGGTCAG AAATGCAAC AGGTTACAC AACTACATCA CATTATTGGA	4260
TAATATCCAG AGGGAACGTT GGGTAATGCT AGCAGGAGCT GTGCTTGGGA TTGCAACTTC	4320
AATCTGTTG TTTAACACTA TGAATAGGCT CTACTTGAA GAATTTAGAC GTGCCATTTC	4380
TATCAAACGC ATTGCAGGTC TCAGGTTCTT AGAAATCCAT CGCACTTATC TCTTGCTCA	4440
ACTGGGTGTG TTTTACTGG GATTTGTTGC GAGTGTATTT CTTCAGGTAG AGATAGGAGT	4500
TGCTTCTTA GTCTGTTAC TCTTACTGG TCTATCTCTT TTACAGTTAC ATGTCCAAT	4560
GCAGAAAAGAA AACAAAGATGT CCATGCTTGT TTTGAAGGGA GGTTAATATG ATTGAACCTTA	4620
AACAGGTGAG TAAATCTTTT GGAGAACGAG AGTTATTTTC GAATCTTCA ATGACATTG	4680
AGGCTGGAAA AGCTCTATGCC TTAATTGGTT CAAGTGGTAG CGGAAAACAA ACCTTGATGA	4740
ACATGATTGG GAAATTAGAA CCTTATGATG GGACGATTTT TTACCGAGGT AAAGACTTGG	4800
CCAATTATAA ATCAAGTGAT TTTTCCGTC ACGAATTGGG CTACCTCTTC CAGAACTTTG	4860
GCTTAATTGA AAACCAAAGT ATTGAAGAAA ACCTTAAGCT AGGTCTCATT GGTCAAAGT	4920
TGAGTCGGTC GGAACAGCGG TTGAGGCAGA AGCAGGCTTT AGAACAGTC GGCCTGGTTT	4980
ATCTTGACCT AGATAAGCGC ATCTTTGAGT TATCGGGCGG AGAATCGCAA CGGGTTCCCT	5040
TGGCAAAAT TATCTAAAG AATCCACCCCT TTATTCTGGC AGATGACCCA ACAGCTTCAA	5100

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TAGACCCAGC AACCTCTAG TTGATTATGG AGATTTGCT ATCTCTCGA GATGATAATA	5160
GGCTAACAT TATCGCAACA CATAATCCGG CAATTTGGGA GATGGCTGAT GAAGTGTCA	5220
CGATGGATCA TCTGAAATAA AAATCCTTGT TTTTAATTGC ACGATGAGTT ACTGAAATAT	5280
TATCATGAAT CAAGAATTGG AGTTAATTAA GAATTGTA CT TAATTTAGAA TTGTACTTTA	5340
TTAATATTGA GGTAACTTTT TCTTGATAAA GGAAGAAATA ATGGAGAGGA AGTTAGAATG	5400
AAAAAATTCG ACAATTATAT TATTGAGAAG CCTTGCGATT CTAATTCAA TAAACTGCAA	5460
AAAATCTAA TAATTGAAAG TTTGGTAGAT GATATTTGC AATTTCTCT CAGAATCAAT	5520
AATACTGTAG GAGAGATTT CCTCCTACAA CCGTTTAAAG AAAACTAT CTTTATTCCA	5580
TGTTATTTG AGGAAGATAT TGTAAGTC AAAAGATGATG ATAAAGTTGA GTGGAATTTG	5640
TTAGAATTTC AAAAATTAG AGCATTTTG CCTTAGTAAT CTGTGTTGAA GGCTCAAAAC	5700
CTATGGTAA AAAGTAGCTT TGAAAACGTA TTGCCTCCAA AGATTTAGTT AAATAATGAT	5760
TTAACACAAA AAGAAATTAT TGAAAGTTCTG GAAAGATGTT GTTCACTAT TGAGAAAAGG	5820
TGGAAAAC TTGCGATTTT CACAGAGAAA GGAAGAAAAA GTATAGAAAT ATAGTCAAATT	5880
GAAACAGAA CAGGATAAAA GAAACCTTTG TGCCATATT TTCTCCTTTC GCTTTACAAT	5940
TGGATTGAAC ACCTTTATTG TATCGCGTTT GGAGTTTTT TGGTATAACC TTCGACGCAC	6000
ACCCGCATAG CGGGTGTGTTT TTTTGTCTCG CACCTAACCG AGCGAGACAA ACTAATAGTC	6060
ACTTAATCAA AAAACGCACC ATATCAAAAAA CTAAAAAGTT TGATATCATG CGTCATGTCT	6120
TAAACTAATT GACTATACCT TCTATTCAA TGAGCTTTA ACCAATTGAT TGAGCCAATC	6180
CACTCTAAA ACCAAAGAGC AATTCTCGC TTAGCTGACT CTTCTGAATC TGAACCATGT	6240
ACAACATTGG GGATAATCTC ATTCTCTCCA GCAGCTTTG CAAAATCACC TCGAATAGTG	6300
CCTGGTAAAG CTTCTCTCG ACAGAGTTGCA CCCATCATGG TCCGCCAAGT TTGATTACT	6360
TTGGGACCG AAATGACACC CACAAGAACT GGACCTGAAG TCATGAATTC ACGAATCGGT	6420
GGGTAAAAC TCTGACCAAC CAAGTCCTGA TAGTGCTGGT CAATCAACTC TTCTGAAACC	6480
TGTGAACGAA ACTCCAATT TTGCGATTGTA AATCCACGTT GTTCGATGCC CTTAACACT	6540
TCACCCACTA GCCCTCTTT TACACCACCT GGTGATGTA TAAAGAATGT TTGTTCCATA	6600
CCCGTCTCCT TTGTCAGCTT CTTCTTTA TTTTACCAACA TTTCGTTGGAA AAATGGAGAA	6660
AGTTTCAGA AGAGAGAATG AGAGAACCT CGGGTTCTCT CATTCTCTCT TATTCTACTG	6720
TTTCTTCCAC AGTTCAACG GCAGTATCCA CAACTACTTC TGTTGTTCT TCATTTCTT	6780
CTTCCTCTAC TGGAGGATTA AGGTATCTT CTTCGTTGAC AGCATGTGGT TCAAGGTTAC	6840

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GGTAACGGGC CATAACCAGTA CCAGCTGGGA TGATCTTACC GATGATAACA TTTTCTTAA	6900
GTCCAAGGAG ATGGTCTTTC TTACCACGGA TAGCTGCGTC AGTAAGGACA CGAGTTGTT	6960
CCTGGAAGGA AGCCGCTGAC AAGAAACTGT TTGTTTCAAG TGAGGCTTG GTAATTCCA	7020
TAAGGACTGG GCGACCTGTC GCTGGAACTC CACCTGCGAT AAGGACATCT TTGTTGGCAT	7080
CTGTAAAGTC ATTGATATCC ATGAGGGTAC CCATGAGAAG ATCTGTATCA CCTGGATCCA	7140
TGACACGGAC TTTACGGATC ATTTGACGAA CCATTACCTC GATGTTGTTG TCACCGATT	7200
CTACCCCTTG GCTACGGTAA ACTTTTGTA CTTCACCGAG AAGGTACGTT TCAACTGACA	7260
AGACATCAGG AACTGCAAGG AGACGTTTG GTTGGATAGA ACCTTCTGTC AGAGCAGCAC	7320
CACGGCCTAC TTGGCCCCA ACTTCGACAC GCATACGAGC TGTAATGGA ACGACATATT	7380
CACCTCGCC AGTTTCACCC TTAACAAAGA CTTTCTTGGT ACGAGTTGAT GCATCTTCTT	7440
CGATAGCAGT AACTGTCCT TTAACCTCTG TAATAACCGC TTCCCTTTA GGATTGCGGG	7500
CTTCAAAAGAT TTCTGGACA CGAGGAGAC CCTGAGTGAT ATCGGTATTG GAGGCAACCC	7560
CACCTGTGTG GAAGGTACGC ATTGTAAGCT GTGTACCAAGG TTCCCGATA GATTGGCAG	7620
CGATTGTACC AACTGTTCA CCAACTTCAA CCGCATCACC AGTCGCCAAG TTGATACCGT	7680
AACAGTGACG GCAGACACCG TGACGAGTGT TACATGTAAA TACAGAACGG ATAGTCACMT	7740
CTTCCACACC AGCATTGACA ATTTCACGCG CCTTGTCTTC TGTAATCAAT TCATTTGGAC	7800
CAATAATCAC TGCACCAAGTT TCTGGATGTT TAACAGTTTT CCTTGTCTAA CGACCCCTCA	7860
GACGCTCTTC GAGAGACTCG ATCATCTCTT TTCCCTCTGC GATAGAACGG ATCAAGAGAC	7920
CACGGTCAGT TCCACAGTCG TCCTCACGGA TGATAACGTC TTGGGCAACG TCGACCAAAC	7980
GACGAGTCAA GTAACCTGAG TCGGCTGTCT TAAGGGCCGT ATCGGTCTATA CCTTTACGAG	8040
CACCGTGAGT TGAGAAGAAC ATTTCCAATA CCGACAAACC TTCGCGGAAG TTTGAAAGGA	8100
TTGGCAATT CATGATACGT CCATTCCGGAG CAGCCATCAG ACCACGCTAA CGGGCAAGCT	8160
GTGAGAAGTT TGAGATGTTA CCACGGGCTC CAGAGTCCAT CATCATAACG ATTGGGTCT	8220
TAGGATCTTG GTTAGCAATC AAGCGTTTCTC CAAGTTTTTC ACAGGGCAGCA CGCCATTCA	8280
CTGTAACAGC ATTGTAACGC TCGTCGTCTG TGATCATACC ACCACGGAAT TGTTGGTGA	8340
TTTGTTCGAC ACGTTTGTGT GATTCTTCAA TGATTTCAGC CTTGTCTATCA ACGACTGGGA	8400
TATCGGCAAT ACCCACTGTC AATCCTGCAA GAGTTGAGTG GTGGTAACCG AGGTTCTTCA	8460
TGGCGTCAAG TAGGGCAGAA GTTTCTGTCG TACGGAAACG TTTGAAGATT TCAGCGATGA	8520
TATTTCAGG GTTTTCTTC TTGAATGGAG GGTTGAGCTC AAGATTGCTG ATAGCTTCCCT	8580
TGATATCTCC ACCAAGTGCG AAGAAGTATT TAGCTGGAAC ACCTTCTGTC AAGTTGGCAT	8640

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TGTTTGGTTC TTGCAAGTAT GGTAGCCCC CTGGCATGAT ATCGTTGAAG AGAATTTCAC	8700
CAACTGTTGT AAGCAAGACC TTATGTCCTT GCTCTCTGT CCAAGGCTTG TTGAGGCTGT	8760
CTGTTGCGAT ACCAACACGT GAGTGGAGGT GAACATAACC ATTGCGGTAA GCCATAACCG	8820
CTTCGTCACG GTCTTGAAG ACCATTCCCTT CACCTTCGCG ACCAGCTTCT TCCATGGTCA	8880
AGTAGTAGTT ACCCAAAACC ATGTCCTGAG ATGGAGTAAC TACCGGTTTC CCATCTTCG	8940
GGTTCAAGAT GTGCTCAGCA GCTAGCATGA GGATACGAGC TTCTGCTTGT GCTTCTTCTG	9000
AAAGTGGTAC GTGGATGGCC ATTTGGTCCC CGTCAAAGTC AGCATTGTAG GCTTCACAGA	9060
CAAGTGGTG CAAGCGAAGA GCCTTACCAT CAATCAAGAC TGGCTCGAAG GCTTGGATAC	9120
CCAAACGGTG AAGGGTCGGT GCGCGGTTCA AAAGCACTGG GTGTTCTTA ATCACTTCTT	9180
CAAGGATATC CCAGATACGC TCATCTCCGC GTTCCACCAA GCGTTAGCT GCTTGACGT	9240
TTTGCACGAT ATCACGGCA ACGATTTCAC GCATGACAAA TGTTTAAAG AGTTCAATCG	9300
CCATTTACAG CGGCACACCA CATTGGTACA TCTTAAGAGT TGGACAAACG GCGATAACTG	9360
AACGTCTGA GAAGTCAACA CGTTTACCGA GCAAGTTTG ACGGAAGCGT CCTTGTTAC	9420
CTTTAAGCAT GTGGCTCAAT GATTTCAATG GACGGCTACC TGGTCTGTG ATTGGACGAC	9480
CACGACGACC ATTGTCAATC AAAGCGTCAA CTGCTTCTTG AAGCATAACGC TTCTCATTTC	9540
GAACGATGAT ACCTGGTCA TTTAACTCAA GCAAACGAGC CAAACGGTTG TTACGGTTGA	9600
TAACACGGCG GTAAAGGTCA TTCAAGTCAG ATGAGGCAAA ACGGCCACCA TCCAACGTCA	9660
ACATTGGACG AAGATCTGGT GGGATAACCG GAAGGATGTT AAGAATCATC CATTCAAGGTT	9720
TGTTTCCAGA CTTGTAAAAG GCATCCAAAA CATCCAAACG ACGGATGGCT TTGACACGGCT	9780
TTTGTCCAGT AGCTGTTTC AATTCTCTT TGAGTTCAAGC AATTCTTTT TCAAGATCTA	9840
CTTGCTTCAA AAGGTCTGG ATGGCTTCCG CACCCATCTT GGCAACAAAT GAACCATAAC	9900
CATAATTCACG CAAGCGCTCT CGGTATTCCGC GCTCTGTAT GATAGACTTG TGCTCAAGTG	9960
GTGTATCCTT AGGATCAATC ACCACATAAG CCGCAAAGTA GATAACTTCC TCGAGGGCAC	10020
GAGGGCTCAT ATCAAGGGTC AAGCCCACAC GGCTTGAAT CCCCTTGAAG TACAGATGTC	10080
GAGATAACAGG AGCTTCAAT TCGATATGTC CCATACGCTC ACGACGAACG TTCTGACGCG	10140
TTACTTCAAC CCCACAGCGG TCACAAACAA TTCTCTGTG ACGAATGCGT TTGTACTTAC	10200
CACAAGCACA TTCCCAGTCT TTTGTAGGAC CAAAGATCAC TTCATCAAAG AGTCCTTCAC	10260
GTTCAGGTTT CAAGGTACGA TAATTGATTG TTTCAGGTTT TTGACTTCT CCATAAGACC	10320
ATGAACGGAC TTTACTTGGA GAAAGCTAGGG TGATTTGCAT ACTTTAAAA CGATTTACAT	10380

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CAACCACTAT TTCTTCCCTT TCTATTCTAA GTGAACTGCT TATTCTTGTT CAGCAGCTTC	10440
TTCTGTTGCT TCCGCTTTG TTGCTTCTC AGCTTCTCA GCTTCAAAGG CTGCTTAGC	10500
CTCTGGGCT GCTTTTCGC GGGCTTTTC AAGGTCATCT ACCTGGATGA CATCTCGTC	10560
CATTCTTCA TCCAAGTCGC GAAGTCCAC TTCTGGTCA TCTTCGTCTA GGACACGCAT	10620
GTCAAGACCA AGAGATTGCA ATTCTTGAC AAGAACCTCG AAGGATTCTG GAACACCTGG	10680
TTTTGGAATT GGTTGCCTT TTGTAATAGC TTCATAGGCT TTCAAACGTC CGTTGATATC	10740
GTCCGACTTG TAAGTCAAGA TTCTTGAAAG GACATTTGAC GCACCGTAGG CTTCAAGAGC	10800
CCAAA CCTCC ATCTCACCGA AACGTTGTCC ACCAAACTGA GCCTTACCTC CGAGTGGTTG	10860
TTGGGTAACA GTTGAGTATG GTCCGACTGA ACACGGCTGC AATTTATCAT CAACCATGTG	10920
GTGGAGTTG ATCATGTACA TGACTCCGAC AGAAACACGG TTATCAAACG GTTCACCAGT	10980
ACGTCCATCG TAAAGGATCG TTTGGCATE GCTATCCATA CCTGCTTCTT TAACAGTTGA	11040
CCAAA GATCT TCAGAACTTG CTCCATCAA GACTGGTGTAA GCGATGTGAA TACCAAGAGT	11100
ACGAGCTGCC ATACCAAGGT GAAGCTCCAT AACCTGACCG ATATTCAAC GTGATGGTAC	11160
CCCAAGTGGG TTCAACATGA TGTGACTGG AGTTCCGTCT GGAAGGTAAG GCATGTCTTC	11220
TACAGGAACG ATACGAGAGA CAACCCCTT GTTTCGGTGA CGTCCGGCCA TTTTATCTCC	11280
GACCTTAATC TTACGTTTT GAGCGATGTAA AACACGAACC AACATGTTAA CACCTGATTG	11340
CAACTCATCT CCATTTACAC GTGTAAGAT CTTAACATCA CGAACGGACAC CATCGGCACC	11400
GTGTGGTACA CGAAGAGAG TATCACGCAC TTCACGAGAC TTGCTCTCAA AGATAGCGTG	11460
CAAGAGACGT TCTTCAGCTG AAAGATCTTT CTCACCCCTTA GGTGTTACTT TACCTACAAG	11520
AATATCACCT TCTTTAACCT CAGCACCAAT ACGGATAATC CCCATTTCGT CAAGGTCTTT	11580
GAGGGCATCT TCACCAACGT TTGGAATTTC CGGAGTGATT TCTTCAGGCC CAAGCTTGT	11640
ATCGCGCTT TCTGATTGTT ATTCTTCAGA GTGAAACAGAT GTGTAGACAT CGTCCTTCAC	11700
CAAGCGTCTG CTCATGATAA CGGCATCCTC GAAGTTGTAA CCTTCCCAG TCATGTAGGC	11760
AACGATTGGG TTTTGTCCAA CGGCCATTTC TCCATTTCGTT ATAGAAGGTC CGTCAGCGAT	11820
GAAATCGCT TTTTCAACGA CATCACCAAC TTTTACGAGA GTGCGTTGGT TGTAAGCAGT	11880
ACCTGAGTTT GAACGACGGA ATTTTGGAT GTGGTAAACA TCCAATGAAC CATCTTCACG	11940
ACGAACCTCT ACCTTGTCAG CATCTGCCTA AGTAACCTTA CCATCTACACT GAGCAATCAC	12000
AGCCGCACCA GAATCGTGGG CTGCTTGGTA TTCCATACCA GTACCAACGT AAGGTGCTG	12060
AGGATTAATC AATGGCACAG CCTGACGTTG CATATTGGCT CCCATGAGGG CACGGTTGGA	12120
GTCACTCGTT TCCAAGAAAG GAATACATGC TGTCGCAACG GCAACTACCT GTTTGGTGA	12180

813

AACGTCCATG TAGTCAACAA TATTAGCTGG ATACTCTTGG TTGACCCCTT GGTGACGTCC	12240
CATGACAATC TTCTCAGCAA AGGTCCATC TTCATTCAA CGAGAGTTAG CCTGAGCTAC	12300
AGTATATTCA TCTTCTTCAT CAGCTGTCAA CCAAACAATT TCGTTCGTGA CAACACCTGT	12360
TTCACGGTCA ACCTTACGGT ATGGTGTTG AACAAAACCA TATTGTTCA AGTGTCCATA	12420
AGATGACAAG TTATTGATCA ACCCGATGTT AGGTCCCTCA CGTGTCTGA TTGGACACAT	12480
ACGACCATAG TGAGTGTAGT GCACGTCACG TACTTCATAT CCAGCACGGT CACGAGTCAA	12540
ACCACCAAGG CCTAAGGCTG ACAAACGGCG TTGTGAGAC AACTCAGAA GCGGGTTGTG	12600
TTGGTCCATG AACTGTGACA ACTGTGATGA ACCAAAGAAT TCTTTAACTG CAGCTGTTAC	12660
AGGACGGATA TTGATAATTT GTTGTGGTGT CAAGACTTCA TTGTCCCTGAA CAGACATACG	12720
TTCACGGACA TTACGTTCCA TACGAGAAAG TCCCACCGT ACTTGGTTGG CAAGCAATT	12780
ACCAACCGCA CGGATACGAC GATTCCAAG GTGGTCGATA TCATCTACAC GGCAAGTCC	12840
TTCAGCCAAG TTGAGGAAGT AGCTCATCTC AGCAAGGATA TCTGCAGGAG TCACCGTACG	12900
AACCTGTCA TCTGGGTTAG CATTACCAAT GATCGTTACG ACCCGATCTG GATCAGTTGG	12960
AGCAATAACC TTGAATTTT GAAGAACAAAC AGGCTCAGTC ACAACGGCTG CATCGTTGG	13020
GATGTAGACA ATCTGTTCA AGTCGCCATC CAAATGGCTT TCAATGCTTT CAATCACGCT	13080
ACGAGTCATA ATCGTACCAAG CTTCTACCAA GATTCTCCA GTTTCAGGGT CTACCAATGG	13140
CTCTGCAATG GTTGGTTGA GCAACAGTGT TTTAACATTG AGTTTTTTAT TGATTTGTA	13200
ACGACCAACT GCTGCCAAGT CATAACGACG TGGCTCAAAG AAGCGAGCTA CAAGCAAGCT	13260
ACGTGAGCTT TCAGCCGTCT TAGGCTCACC TGGACGAAGG CGTTCGTAAA TTTCTTCAA	13320
GGCTTCGTCT GTACGAGAGT CCATTGGATT CTTGTGGATA TCTTTTTCAA CAGTGTGCG	13380
AACCAATTG CTGTACCAA AGATATCAA GATTCATCA TCACCTGAGA AACCAAGAGC	13440
ACGAACCAAG GTTGAAATG GAATCTTACG AGTACGGTCG ATACGAGTGT AGGTGATATC	13500
TTTGAGTCG CTTCAAGTT CCAACCAAGC TCCACGGTTA GGGATAACAG TTGAACCATA	13560
GCCCCACCTTA CCATTTTGT CTACTTTGTC GTTAAAGTAA ACACCTGGTG AGCGGACCAA	13620
CTGAGAAACG ATAATACGTT CACCACATT GATGATGAAA GTACCCATT CTGTCATGAT	13680
TGGGAAATCA CCAAAGAAA CTTCTGGGT CTTGATTTCG CTTGTTCTT TATTGATCAA	13740
ACGGAAAGTT ACAAAATTG GTGCTGAGTA GCTAGCATCG TGGATACGAG CTTCTTCTAG	13800
CGTATATTGTT GGTCCTTGA TTTCATATCC AACAAATTCC AACTCCATTG TGTCTGTGAA	13860
GTTGAAATT GGCAATACAT CTTCAAACAC TTCCCTTAAGA CCGTGGTCTA GGAAAGCTT	13920

814

GAATGAGTCA	GTTTGAATT	CAATCAAATT	TGGTAAGTCA	AGAAACTTCTT	TGATTCTTGA	13980
AAAAC TACGA	CGGGTACGAT	GTTTCCCGTA	TTGAACGTCA	TGTCCTGCCA	AGATGATTCT	14040
CCTTTGTAAA	TAAGTTCCAA	GCCTTGTCAA	TCAGGCTTT	CTAATCGTCA	TATGGTTGTA	14100
AACCCCTTAT	CACCGTGTCC	TCTTGACGAA	TTTCAGAAT	CTTTAAGCCT	CTGTTACAAA	14160
TGCTCAAAAT	CTTGAAAAAA	AGCACAAAAA	GAGCAGCTAA	ATCTGACTTT	TTCAGAAGAT	14220
TTAAC TGCTG	TGAGCCTTGT	CTGGACAATA	TTTCAGACAA	AACTACGAC	AAATGATTAC	14280
CCATATTATA	CCCTATTTAG	CTAGATTTT	CAAGGGTTT	CGTAGGTTT	TTGGTAAATT	14340
TTTTCCCATA	GAAAACTTGG	CATCACATTC	GAATCACGCT	ATGTCACAAA	AAACTGAAAA	14400
AACTATMGC	TGAAAATCAT	TTTCAAGGTA	TAATAATAAA	CGTTAAGGCG	GTATAGCCAA	14460
GTGGTAAGGC	ACGGCTCTGC	AAAAGCTTGA	TCGTCGGTTC	AAATCCGTCT	ACCGCCTTCT	14520
ATAACTTGAT	TTATCAGGTT	TCAAATGAAC	AGAAAAGCCA	ATTGAAAGGG	CTTTTTTAT	14580
TTTCCCTCGA	ATAAAATACGT	ATAACTTTAA	AAACTTTGG	AGCGAGTTG	TGGCAGAGTT	14640
CTTTCCATGG	CATAATTCCC	TTTGAAATC	AG			14672

(2) INFORMATION FOR SEQ ID NO: 112:

- (i) SEQUENCE CHARACTERISTICS:
- (A) LENGTH: 7902 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 112:

AGGAGACTAT	TCAAGCCAA	ATTgAGTAGC	CCAGCAAAGA	CTGTATAGAC	TGTGATACTG	60
TTTTCATAGC	CATTGGTAAA	GAGAATTGG	GAACCAAGAA	TGGTATCTAA	GGCCAGGATA	120
ATCGTACGAA	AAGCGAAGAG	AGAGGTCAAG	ATGCCGCCTC	CGATATATTT	TTCACTACCG	180
TAAAGTAGGA	TGGCATTGG	TCCTAAAACC	ATGAGTCAA	AACTCAGTGG	AATGATAAAG	240
AAAGTAAAGA	TTCGACTACC	TCTATTAACC	AGAGAAACAT	AGGCTTCTTT	GTCTCCTTTC	300
CCCAGATAGT	AACTGAGACG	AGGCACACTC	ACTCCAATTG	CACCTGTTAC	AAACCCAGCT	360
ATAACGGTCA	CAATTGCTG	AGCTATGGTA	TAGTAACTAA	CGTTGACATC	AATCCCTGTT	420
TTAACCGAGGA	AGAGGGCAGTC	AAAAAAAGTG	AGAGGCATAT	TGGCATGGC	AAAGACTAAC	480
ATGGCTGTCA	GAGGGAGAAA	GAGTGGTTA	AAATCACTTA	GGTGAATT	AAACAGTTG	540
ATGTCCTTT	TAATCCAAA	ATAACTAAC	AGGTAGTTAA	TCAGCGTCGA	AAAACATC	600
ACAAGTGTAT	AGACAACAAT	ATCGTGTCA	TTTTAACAA	ATAAGAAAAT	AGAGACCAGC	660

815

ATCAGGATAC GGATGAAGGC AGTTTGTAA AAGAGAAAAC TGTAATTTC CAGAGCTTC	720
TTGACCCATT CGATTGAAAA AATCTGGCA ATGAGTTGAA TCCCCATAAC AAGGTAGACC	780
TTTTTGACGA TTGGATTATC AGTAAAGAAG AGAGGATAGG CTAGGATATA GACAGCAGTG	840
GTCAAATCG TACAAGCGAT GCACAAATAA AAAAGACTAG AAAAGGTTCT GTTAAGATCT	900
TTTTTGTAT CCTTGACATT ACTGATGCC CTTAAACCGT AGTTATAGAC ACCATAAGTT	960
GCAAAGGGCA AGAAAATGA CAAAATAGTG TCGACTGAGT TGAAGTAACC ATAGTCAGTT	1020
CGGTCCAAGA CACGCCGCAC ATAGGTTCCA GTTAGGATGG GAAAATAAT ATTCAAGACA	1080
CGAATTCCCA TGTAAGATAG AGCATTTAAT TTTATACTTT TCATTCAATT TACCTCGTTT	1140
TTCATTATAT CATAAAAGTTA GCTAATAAGA AATGAAGGGC AGTAAGTCAA GTAATCAGTT	1200
TGAAGTTCA AATCTTAAGT TTTAAGTTT CTTAAGGAA AGTATATTAT TCTGAAGGAC	1260
TCTAAAATTT CGCAGCCATT TATTAGTAAT TGCTACAGAA TTCCTAGTCA TTACTAGAAA	1320
TGGACTAGTT TCTTTGAATA ATAGAACTGC ATAATTCTCC TATTCTAGAA GGGGAGGACC	1380
AGTATTCTT TTATGATAGG ACTAGATTGT GGTATAATAG AGAGAATAAG TTTTTTTAGT	1440
AAGACAAAGG AGAAAATAGA TGATTATGC AGGAATTCTT GCCGGTGGAA CTGGCACACG	1500
CATGGGGATC AGTAACCTGC CAAAACAATT TTTAGAGCTA GGTGATCGAC CTATTTGAT	1560
TCATACAAATT GAAAATTG TCTGGAGCC AAGTATTGAA AAAATTGAG TTGGTGTCA	1620
TGGAGACTGG GTTTCTCATG CAGAAGATCT TGTAGATAAA TATCTTCCTC TTTATAAGGA	1680
ACGTATCATC ATTACAAAGG GTGGTCTGA CCGCAATACA AGTATTAAGA ACATCATTGA	1740
AGCCATTGAT GCTTATCGTC CGCTTACTCC AGAGGATATC GTTGTACCC ACGATTCTGT	1800
TCGTCCATT ATTACACTTC GCATGATTCA GGACAATATC CAACTTGCCTT AAAATCATGA	1860
CGCAGTGGAC ACAGTGGTAG AAGCGGTTGA TACTATCGTT GAAAGTACCA ATGGTCAATT	1920
TATTACAGAT ATTCCAATC GTGCTCACCT TTATCAAGGA CAAACACCTC AAACATTCCG	1980
TTGCAAGGAC TTCATGGACC TTTATGGATC TCTTTCTGAT GAAGAGAAGG AAATCTTGAC	2040
AGATGCATGT AAAATCTTG TGATCAAAGG AAAAGATGTG GCTTTGGCCA AACGTGAATA	2100
CTCAAATCTG AAGATTACAA CGTAACAGA TTTGAAGATT GCAAAAAGTA TGATTGAGAA	2160
AGACTAGTAA AATGATTAAT CAAATTATC AACTAACTAA GCCTAAGTTT ATCAATGTCA	2220
AATATCAGGA AGAGGCTATT GACCAAGAGA ATCATATCCT TATCCGTCCC AACTACATGG	2280
CTGTCTGTCA TGGCGATCAG CGTTACTATC AGGGAAAACG TGATCCCAAG ATTTTGAAATA	2340
AAAAGCTTCC AATGGCAATG ATTACACGAGT CATGTGGAAC CGTCATTCT GACCCGACCG	2400

816

GAACCTACGA GGTTGGTC	AAAGTTGTCA TGATTCCAA	TCAGTCTCCT ATGCAGAGTG	2460
ATGAAGAAC	CTATGAAAAC TACATGACAG	GGACCCATT CTTGTCAGT GGATTTGATG	2520
GCTTTATGAG AGAGTTGTT	TCTCTCCCTA AAGATCGTGT	GGTGGCTTAT GATGCTATTG	2580
AAGATAACGGT	TGCAGCCATT ACAGAGTTG	TCAGTGTGGG CATGCACGCT ATGAATCGTC	2640
TATTGACTCT	TGCTCATAGC AAGCGGGAGC	GGATGCCGT TATTGGAGAT GGAAGTTAG	2700
CTTTTGTGGT	TGCCAATATT ATCAACTATA	CTTGCCAGA AGCAGAGATT GTGGTTATTG	2760
GTCGTCATTG	GGAAAAGTTG GAACTCTCT	CATTTGCCAA AGAATGCTAT ATTACGGATA	2820
ATATTCTGTA	AGATTTGCC	TTTGACCATG CTTTGAAATG TTGTGGTGGT GATGGTACTG	2880
GACCAGCTAT	TAATGACTTG ATTCGCTACA	TTCGTCCTCA GGGAACGATT CTCATGATGG	2940
GAGTTAGCGA	ATATAAAGTC AATCTCAATA	CTCGCGATGC CTTAGAAAAG GGCTTGATTT	3000
TGGTTGGGTC	ATCTCGTTCT	GGTCGCATTG ATTTGAAAAA TGCTATCAA ATGATGGAAG	3060
TCAAGAAATT	TGCCAATCGT CTTAAAATA	TCCTTTATCT AGAAGAACCT GTAAGAGAAA	3120
TTAAAGATAT	TCATCGTGT	TTTGCAACCG ATTAAACAC AGCCTTTAAA ACAGTGTAA	3180
AGTGGGAAGT	ATAAGTACTG GAGGTTAATT	GTGGAGAAAA TCATTAAAGA AAAAATTCT	3240
TCCTTACTTA	GTCAAGAAGA GGAAGTCCTC	AGTGTGAAAC AACTGGGTGG AATGACCAAT	3300
CAAAACTATT	TGGCCAAAAC AACAAATAAG	CAATACATTG TAAATTCTT TGGTAAAGGG	3360
ACAGAAAAG	TTATCAATCG ACAAGATGAA	AACTACAATC TTGAACACT AAAGGATTTA	3420
GGCTTAGATG	TAAAAAATT	TCTTTTGAT ATTGAAGCTG GTATCAAAGT AAATGAGTAT	3480
ATCGAATCTG	CGATTACGCT	TGATTCAACG TCAATCAAGA CCAAGTCGA CAAAATTACT	3540
CCAATATTAC	AAACTATTCA	TACCTCTGCT AAGGAATTAA GAGGAGAATT TGCTCCTTT	3600
GAAGAAATCA	AAAAATACGA	ATCCTTGATT GAAGAACAAA TTCTTATGC CAACTATGAA	3660
TCTGTTAGAA	ATGCAGTCTT	CTCCTTAGAG AAAAGACTGG CTGACTTAGG TGTTGACAGA	3720
AAATCTGTC	ATATCGATT	GGTCCTGAA AACTTTATCG AATCACCTCA AGGACGACTT	3780
TATTTGATTG	ACTGGAAATA	TTCATCAATG AATGATCCAA TGTGGGATTT GGCTGCCCTC	3840
TTTTTAGAGT	CTGAATTCA	TTCCAAGAG GAAGAACATT TCTTATCTCA CTATGAGAGT	3900
GACCAACAC	CGGTTCTCA	TGAAAAGATT GCTATTATA AAATTTACA AGATACTATT	3960
TGGAGTCTAT	GGACTGTCTA	TAAGGAAGAG CAAGGTGAAG ATTTGGTGA CTATGGTGTG	4020
AATCGTTACC	AAAGAGCTAT	TAAGGTTTG GCTTCTTATG GAGGTTCAGA TGAAAAGTAA	4080
AAACGGAGTT	CCTTTGGCC	TTCTCTCAGG TATTTCTGG GGCTTGGTC TAACGGTTAG	4140
TGCTTATATC	TTTCGATT	TTACAGATT GTCACCCATT GTGGTGGCTG CAACTCATGA	4200

817

TTTTTTGAGC ATCTTATCT TACTAGCTT TCTCTGGTA AAAGAAGGGA AAGTTGCCCT	4260
CTCAATTTC TTAAATATTC GCAATGTCAG TGTTATCATC GGAGCCTTGC TAGCAGGCC	4320
TATCGGTATG CAGGCCAATC TTTATGCAGT TAAGTATATC GGAAGTCTT TAGCTTCATC	4380
TGTATCGGCT ATTACCCCTG CGATTTCACT TCTATTGGCT TTCTCTTTT TGAAGCACAA	4440
GATTCGAAA AATACTGTAT TTGGGATTGT CTTGATTATT GGAGGGATTA TTGCTCAGAC	4500
CTATAAGGTT GAACAGGTTA ATTCTTCTA CATTGGGATT CTTTGTGCTT TGGTTTGTGC	4560
TATTGCATGG GGAAGTGAGA GTGTTCTTAG CTCTTTGCC ATGGAAAGTG AATTGAGTGA	4620
AATCGAAGCC CTCTTAATCC GTCAAGTAAC TTCGTTCTTG TCCTATCTTG TGATTGTGCT	4680
CTTCTCTCAT CAGTCATTTA CTGCAGTAGC CAATGGACAA TTGCTAGGTC TCATGATTGT	4740
TTTTGCAGCC TTTGATATGA TTTCCTACTT GGCTTATTAT ATCGCTATCA ATCGCTTGCA	4800
ACCAGCCAAG GCTACAGGCT TGAACGTGAG CTATGTAGTA TGGACGGCT TGTTTGAGT	4860
TGTTTCTTG GGTGCACCGC TAGATATGCT GACCATTATG ACGTCACCTTG TCGTCATTGC	4920
TGGAGTTAT ATTATTATTA AAGAATAAAG GAGATTCTGT TGAAAGCCAT TATCTTAGCA	4980
GCGGGATTGG GAACTCGCTT GCGCTCTATG ACTGAAAATA CCCCTAAAGC CTTGGTTCA	5040
GTAAATCAA AACCTTGAT TGAGTACCAA ATTGAGTTTC TCAAAGAAAA AGGAATCAAT	5100
GACATCATCA TCATPTGTTGG TTATCTAAA GAACAATTG CATTACTGAA AGAGAAATAC	5160
GGTGTTCGTC TCGTTTCAA TGATAAATAC GCTGACTACA ATAACTTTA CTCTCTCTAT	5220
CTTGAAAAG AAGAATTGGC CAACAGCTAT GTTATTGATG CTGACAATTAA TCTCTTTAAA	5280
AATATGTTCC GCAATGATTT GACACGTTCG ACTTATTTA GTGTTATCG TGAAGATTGT	5340
ACCAACGAAT GGTTCTGGT TTATGGAGAT GACTACAAGG TTCAAGACAT TATTGTTGAT	5400
AGCAAGGCAG GTCCGATCCT TAGTGGTGT TAATGGTGTG ATGCTCCAAC TGCAGAAAAG	5460
ATTGTCAGCT TTATCGACAA GGCTTATGTA AGTGGTGAAT TTGTTGATCT CTATTGGGAC	5520
AATATGGTTA AGGATAATAT CAAAGAGCTA GATGTCTATG TTGAAGAATT AGAAGGCAAT	5580
AGCATTATG AGATCGATAG TGCTCAAGAC TATCGTAAAT TAGAAGAAAT TCTTAAAAC	5640
GAAAATTAAA GATTCCAACA TCTGACAAAA TAGTCGGATG TTTTTGATT TTTTACGAAC	5700
TTTTACGAAT AGATAGATGA GTAGAAAAAG AAATGGAGTT ATTTATGAAA ATCACAAACT	5760
ATGAAATCTA TAAGTTAAA AAATCAGGTT TGACCAATCA ACAGATTTG AAAGTGTAG	5820
AATACGGTGA AAATGTTGAT CAGGAGCTT TGTTGGGTGA TATTGCAAGAT ATCTCAGGTT	5880
GCCGTAATCC AGCCGTTTT ATGGAACGTT ATTTTCAGAT AGACGATGCG CATTGTCGA	5940

818	
AAGAGTTCA AAAATTCCA TCTTCTCTA TTTAGATGA CTGTTATCCT TGGGATTTGA	6000
GTGAAATATA TGATGCGCCT GTACTTTAT TTTACAAGGG AAATCTTGAC CTCCTGAAAT	6060
TCCCGAAGGT AGCGGTCGTG GGCAGTCGTG CTTGTAGCAA ACAGGGGACT AAGTCAGTTG	6120
AAAAAGTCAT TCAAGGCTTG GAAAATGAAC TGGTTATTGT CAGTGGTCTG GCCAAGGGCA	6180
TTGACACAGC AGCTCATATG GCAGCTCTTC AGAATGGCGG AAAAACCAATT GCAGTGATTG	6240
GAACAGGACT GGATGTGTTT TATCCTAAAG CCAATAAACG CTTGCAAGAC TACATCGGCA	6300
ATGACCACATCT GGTTCTAAAGT GAATATGGAC CTGGTGAACA ACCTCTGAAA TTTCATTTTC	6360
CTGCCCGTAA TCGCATCATT GCTGGACTTT GTCGTGGTGT GATTGTAGCA GAGGCTAAGA	6420
TGGCGTTCAGG TAGTCTCATT ACGTGTGAGC GAGCAATGGA AGAAGGACGC GATGTCCTTG	6480
CTATTCCCTGG TAGCATTTTA GATGGACTAT CAGACGGTTG CCATCATTG ATTCAAGAAG	6540
GAGCAAAATT GGTCACCAAGT GGGCAAGATG TTCTTGCAGA ATTTGAATT TAAAATGAC	6600
CTAAGCTAGA ATTCTAAGAA AAAATCAATT TTAAGAGAAA ATGAACCCAA CATTCCATA	6660
ATAAAACGCA TATTAGCAAG TTTTTAACAC TTGATAATAT GCGTTTTTC TAAGTGGATT	6720
AGTAGAGTAG AGGATTTTC TCATATAATA CTCTTCGAAA ATCTCTCAA ACTACGTCAG	6780
CTTCCATCTG CAAACCTCAAA ACAGTATTTT GAGCgACTtC GTCAGTCTTA TCTACAAACCT	6840
CAAAGCAGTG CTTTGAGCAA CCTGTGGCTA GCTTCCTAGT TTGCGCTTTG ATTTCATTTG	6900
AGTATAAGGG AAAGTATAGT GAATTGAAAT AAGATGTGAA CAACTCTATC AGGAAAGTCA	6960
AATTAATTAA TAGAAATATT TTACGAGCCA AGGTGTACTG TTATAGATTC AATTACACTA	7020
TAATTTAGTG TAATTGAGAA AGGAGAAATG ATTGTGATTG ATGTTGGCTA GGTTATGTT	7080
AATGATTCCT ACCGCTCTCAA ATCTTGTCAAG TAAGGAAAAA TAAATTCTTC AAAAGTAGAG	7140
ATTACAAGGC TTGTTAAGA AAGAATTCAA AGACCTTGAC AAATAAAAAT AAAATGGTTA	7200
TTATAAAAAA TGGCTGAAA TAGATGATGA TACTTTCGA AAATCTCTTC AAATACGTCA	7260
GCTCAGCTTT GCCTTGCTGT GTTTGAGCA AGCTACGGTT AGCTTCCGAG TTTGATTTTC	7320
ATTTACTAGA AATGAAACTG ATGAGAGATA TCAGTAGACA TTTGAGTCAG GATATTATGG	7380
AAAATGATAA AAAGAGCTCG TGAGATTGGC ATATCAGACT ACTAAAGTAT TGAGTTGTT	7440
AGGATTTAG CGACTAGTTA GCTGGAAAG GAAGATATTG GTGACAAATA ATAAACTGTA	7500
TTCGTTGATA GAATTTAGAA ATAAAATATA TGAAGAATTA GAACTTCCA GAAAGTGT	7560
AGCGATTTA CTATGTGCCA TGCTTATCGC CTCTATCGGA TTAAATATGG ATTCGACTCC	7620
CGTGATTATT GGAGCCATGT TAATCTCTCC TTTGATGACA CCTATTCTGG GAGTGGGGCT	7680
CTCTCTAGCT ATATTTGATT TTAAATGTT AAGAAAATCT TTAAATATAT TAGCTATTCA	7740

819

AATTCTTGCC AGTCTAATAG CTTCAACACT TTATTTTAT CTTTCTCCC AATCGTATGC	7800
TAGTTGGAG ATTGTTGCTA GAACCTCTCC GACTATTTGG GATGTTCTCA TTGCTTTGT	7860
AGGAGGGATA GCAGGTATCA TTGGTGCTAG GAAAAAAGAG AC	7902

(2) INFORMATION FOR SEQ ID NO: 113:

- (i) SEQUENCE CHARACTERISTICS:
- (A) LENGTH: 18627 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 113:

GAAGTTGAAA TGGCCAGCTG ATGAGCAATA TCGGTCTAG AAATCTTCTC AATCAACTTT	60
TGCGCAATT TTTGGGTGAT AATACGAGGA ATTTGGTGAT TTTTCTTGAC GATAGAAGTT	120
TCAGCGACCA TCATTTTGA ACAGTGATAG CACTTGAAAC GACGTTTCT AAGTAGAATT	180
CTAGTAGGCA TACCAAGTTGT CTCAAGGTA GGAAATCTTAG ACGGTTTTG AAAGTCATAT	240
TTCTTCAATT GGTTCCGCA CTCAGGGCAA GATGGGGCGT CGTAGTCCAG TTTGGCGATG	300
ATTTCCCTGT GTGTATCTTT ATTGATGATG TCTAAATCT GGATATTAGG GTCTTTAATG	360
TCTAGTAATT TTGTGATAAA ATGTAATTGT TCCATATGAA TCTTTCTAAT GAGTTGTTTC	420
GTCGCTTTTC ATTATAGGTC ATATGGGACT TTTTTCTAC AATAAAATAG GCTCCATAAT	480
ATCTATAAGG GATTTACCCA CTACAATAT TATAGAGCCA AAAATCCTTT GTTTACTAAA	540
CAAGGGATTT TTCTTTGTC TCTGCTCCTT TTTGATATA ATAGTTCTAT GTTAAATCA	600
GAAAAACAAT CACGTTATCA AATGTTAAAT GAAGAATTGT CCTTCCTATT GGAAGGCGAA	660
ACCAATGTTT TGGCTAATCT TTCCAACGCC AGTGTCTCA TAAAATCACG TTTTCTTAAT	720
ACCGTATTTG CAGGTTTTA TTTGTTGAT GGAAAGGAAT TGGTTTTAGG CCCCTTCCAA	780
GGAGGTGTTT CCTGCATCCG TATTGACTA GGCAAGGGTG TTTGTGGTGA GGCAGCTCAC	840
TTTCAGGAAA CTGTTATTGT TGAGATGTG ACGACCTATC TCAACTATAT TTCTTGTGAT	900
AGTCTAGCTA AAAGTGAAT TGTGGTCCCG ATGATGAAGA ATGGTCAGTT ACTTGGAGTT	960
CTGGATCTGG ATTCTTCAGA GATTGAGGAT TACGATGCTA TGGATCGAGA TTATTTGGAA	1020
CAATTGTCG CTATTTGCT TGAAAAGACA GCATGGGACT TTACGATGTT TGAGGAAAAA	1080
TCTTAATGTA TCAAGCACTT TATCGAAAAT ATAGAAGTCA AAACCTCTCC CAGTTAGTTG	1140
GTCAAGAAGT TGTGGCTAAG ACTCTTAAAC AACCGGTGGA GCAAGAGAAA ATAAGTCACG	1200

820	
CTTATCTTT TTCTGGCCT CGTGGAACGG GAAAACCAG TGGTGCCTAA ATCTTGCCA	1260
AGGCTATGAA CTGTCCCAAT CAAGTGGGTG GCGAACCTTG CAATAACTGC TATATTGTC	1320
AAGCAGTGAAC GGACGGTAGT TTAGAAGATG TCATTGAAAT GGATGCGAGCT TCTAATAATG	1380
GGGTAGATGA AATTGCGAA ATTCTGTGATA AATCTACCTA TGCGCCTAGC CTTGCTCGTT	1440
ATAAGGTTTA TATCATAGAT GAGGTCACA TGCTGTCTAC AGGGGCTTTT AATGCCCTCC	1500
TAAAGACGCT GGAAGAACCA ACACAGAACAT TAGTCTTTAT TTTGGCCACT ACTGAATTGC	1560
ACAAGATTCC TGCTACTATT CTATCCCCTG TGCAACGTTT TGAGTTAAA TCAATTAAGA	1620
CACAGGATAT TAAGGAACAT ATTCACTATA TCTTAGAAAA AGAAAATATC AGTTCTGAAC	1680
CAGAGGCTGT GGAAATCATT GCCAGACGGG CGGAAGGTGG AATGCGGGAC GCCTTGCTA	1740
TTTTGGATCA AGCCCTGAGT TTGACACAGG GAAATGAGCT GACGACTGCT ATCTCTGAAG	1800
AAATTACTGG CACCATAGC CTATCAGCCT TGGATGATTA TGTGGCGGCC TTGCTCAAC	1860
AGGATGTTCC CAAAGCTTG TCTTGCTTGA ATCTTCTTT TGACAATGGT AAGAGCATGA	1920
CTCGTTTGTG GACCGATCTT TTGCACTATT TAAGAGACTT GTTAATTGTT CAAACAGGGG	1980
GAGCAAATAC TCATCATAGT TCAGTCTTG TAGAAAATTT GGCACCTCCT CAAAAAAATC	2040
TGTTTGAAAT GATTGCTTA GCAACAGTGA GTTGTAGCAGA TATTAAGTCT AGTTGCAAC	2100
CCAAGATTTA TGCTGAAATG ATGACCGTCC GTTGGCGGA AATCAAGTCC GAACCAGCTC	2160
TATCAGGAGC GGTTGAAAT GAAATTGCTA CGCTGAGACA GGAAGTGTGCG CGTCTCAAAC	2220
AAGAGCTTTC TAATGTAGGT GCGGTTCTA AACAAGTTGC ACCAGCTCCT AGTCGACCAG	2280
CTACGGCCTA AACAGTCTAT CGTGTGATC GCAATAAGT GCAATCTATC TTACAAGAGG	2340
CCGTCGAAAA TCCTGATTTA GCACGTCAAATTTAATCG TTTGCAGAAT GCCTGGGGAG	2400
AGGTAATTGA AAGTCTAGGT GGGCCGGACA AGGCTGTGCT AGTTGGTTCT CAACCGGTTG	2460
CTGCCAATGA ACACCATGCT ATTCTTGCTT TTGAGTCTAA CTTCAATGCT GGTCAAAC	2520
TGAAACGAGA CAATCTCAAT ACCATGTTG GTAATATCCT CAGTCAGGCG GCAGGTTTT	2580
CACCTGAGAT TTAGCTATT TCCATGGAGG AATGGAAAGA AGTTGCGCA GCCTTTTCAG	2640
CCAAAGCCAA ATCTTCTCAA ACTGAAAAAG AAGTAGAAGA AAGCCTGATT CCAGAAGGAT	2700
TTGAATTTTT GGCTGATAAA GTGAAGGTAG AGGAAGACTA AAGAAAGATT TCATGATACA	2760
ATAAGTTTAT GAATAAACAA CAATTATTA TTATGGCGCT GTTACAGCT GCTGAGACCT	2820
ATTTTTCTAA TGAACCTGG ATGACTGGCC GCTATATTAT GGCAGCCTTT TGGGCAATT	2880
TAATCTTTAG AAATTCCGA GTCAAGTTATG TGATGGCAA AATCGTTGAT GTCATCGATC	2940
AGCATTAA TAGGAAAGAC TAGCCCTCAG CTTCCAGACA AAATCAAAGC CTTTTAGGCT	3000

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TTTTTTTGT	ATACTAGAAA	AGTATATT	TAGAATT	TTC GCTCTATT	TGGGGAAATC	3060	
AGACGTTTT	CTAGTAAGTA	CTGTAAAAGT	TTTGAAAAG	AAAGGAAC	TA TCATGTCAGT	3120	
ATTAGAGATC	AAAGATCTTC	ACGTTGAGAT	TGAAGGAAA	CAAATT	TTAA AAGGGGTTAA	3180	
CCTGACCC	CTG AAAACAGGAG	AAATTGCCGC	TATCATGGCA	CCAAATGGTA	CAGGTAAC	3240	
GA	CTCTTCT	GCGC	TATCA	TGGGAAATCC	AAACTATGAA	GTAACTAAAG	3300
GT	TTTGATGGC	GTAAACATCC	TTGAGTTGGA	AGTGGATGAG	CGTGC	GCTA TGGGACTTT	3360
CCTTGCTATG	CAATACCCAT	CAGAAATCCC	TGGAATTACC	AATGCTGAGT	TTCTCGTGC	3420	
CGCTATGAAT	GCGGGTAAAG	AAGATGATGA	GAAGATT	CA	TTCTGAGT TTATTACTAA	3480	
GCTAGATGAA	AAAATGGAAT	TGCTCAACAT	GAAAGAAGAA	ATGGCAGAGC	GTTAC	CTCAA	3540
CGAAGGCTTC	TCTGGTGTG	AGAAAAAACG	CAATGAAATT	CTTC	AACTTT TGATGTTGGA	3600	
GCCAACATTT	GCTCTTTGG	ACGAGATTGA	CTCAGGTCTT	GATATTGAC	CTCTTAAAGT	3660	
TG	TGTCTAAA	GGTGTCAATG	CCATGCGTGG	TGAAGGTTT	GGTGC	TATGA TCATCACTCA	3720
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TGTTGTC	CTCTGGGTC	CAGAATTGGC	TGCGCGTTG	GAACGTGAAG	GATA	CGCAA 3840	
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TGCAAGTATT	GACTGGGCTA	TCGGTGT	CAT GAACGAAGGA	AATGTC	GTTG	CTGATT	TGA 4680
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GATGCTGAGC AAATCAGTCT CAACAATCCG ACTTGTGGGG ATGTCATCAA CCTCTCTGTC	6540

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AAGTTTGATG CAGAGGACCG TTTGGAAGAT ATTGCTTTTC TAAATTCAAGG ATGCACGATT	6600
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CCTTTACAAA AGAACCTTCCA ATGGAATACG CAGTTGAGCT GAACCGCTTG ATTAGCTATG	8280

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AAATGGAGGG ATCAGTTGGA TAAAATTTGA TTTTATACTC TTGAAAATC TCTTCAAACC	8340
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GCTTTTTTA TTATTGACA AACCTATTG TTACTTTCTT AAATAAGTC TGTAAACTCA	10080

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AATTCCCTTG CTAGTGGTA GCCTAGCTTA TCTGGCGATA TTCGGCCTAT TAATCTATCT	16200
CTTCTTTTC AAGTGGATAC GAAAACAGGA AGGACTCTTA TCTGGCTTT TCACCATATT	16260
TGCTGGCTTA CTCTTGATT TTGAGGCCA CTTGGTTGG AAAATATGGTT TGGACAAGTC	16320
CGTTCTAAAA GGGACCATGG CTCAGGTGT GACAGATCTG ACTGGTTTC GAACGACTAG	16380
CTTTGCTGGA GGGGGCTTGA TCGGGGTCGC TCTTTATATT CCAACAGCCT TTCTCTTTTC	16440
AAATATCGGA ACTTACTTTA TTGGTTCTAT CTTGATTTA GTGGGTTCTC TCCTAGTCAG	16500
CCCTTGGTCT GTTACGATA TTGCTGAATT TTTCAGTAGA GGCTTGCCA AATGGTGGGA	16560
AGGGCACGAG CGTCGAAAG AGGAACGCTT TGTCAAACAA GAAGAAAAAG CTCGCCAAA	16620
GGCTGAGAAA GAGGCTAGAT TAGAACAAAGA AGAGACTGAA AAAGCCTTAC TCGATTTGCC	16680
TCCTGTTGAT ATGGAAACGG GTGAAATTCT GACAGAGGAA GCTGTTCAA ATCTTCCACC	16740
TATTCCAGAA GAAAAGTGGG TGGAACCAGA AATCATCCTG CCTCAAGCTG AACCTAAAATT	16800
CCCTGAACAG GAAGATGACT CAGATGACGA AGATGTTCAAG GTGATTTTT CAGCCAAAGA	16860
AGCCCTTGAA TACAAACTTC CAAGCTTACA ACTCTTGCA CCAGATAAAC CAAAGATCA	16920
GTCTAAAGAG AAGAAAATTG TCAGAGAAAA TATCAAAATC TTAGAACGAA CCTTGCTAG	16980
CTTTGGTATT AAGGTAACAG TTGAACGGGC CGAAATTGGG CCATCAGTGA CCAAGTATGA	17040
AGTCAAGCCG GCTGTTGGTG TAAGGGTCAA CCGCATTTC AATCTATCAG ATGACCTCGC	17100
TCTAGCCTTG GCTGCCAAAG ATGTCCGGAT TGAAGCACCA ATCCCTGGGA AATCCCTAAT	17160

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CGGAATTGAA GTGCCCAACT CCGATATTGC CACTGTATCT TTCCGAGAAC TATGGGAACA	17220
ATCGCAAACG AAAGCAGAAA ATTTCTTGG AATTCCCTTA GGGAGGCTG TTAATGGAAC	17280
CGCAAGAGCT TTTGACCTTT CTAAAATGCC CCACCTGCTA GTTCAGGTT CAACGGGTT	17340
AGGGAAAGTCA GTAGCAGTTA ACGGCATTAT TGCTAGCATT CTCATGAAGG CGAGACCAGA	17400
TCAAGTTAAA TTATGATGG TCGATCCAA GATGTTGAG TTATCTGTTT ACAATGATAT	17460
TCCCCCACCTC TTGATTCCAG TCGTGACCAA TCCACGCAAA GCCAGCAAGG CTCTGCAAAA	17520
GGTTGTGGAT GAAATGGAAA ACCGTTATGA ACTCTTGCC AAGGTGGAG TTCGGAATAT	17580
TGCAGGTTTT AATGCCAAGG TAGAAGAGTT CAATTCCCAG TCTGAGTACA AGCAAATTCC	17640
GCTACCATTG ATTGTCGTGA TTGTGGATGA GTTGGCTGAC CTCATGATGG TGGCCAGCAA	17700
GGAAAGTGGAA GATGCTATCA TCCGCTTGG GCAGAAGGCG CGTGCAGCAG GTATCCACAT	17760
GATTCTTGCA ACTCAGCGTC CATCTGTTGA TGTCATCTCT GGTTGATTA AGGCCAATGTT	17820
TCCATCTCGT GTAGCATTTG CGGTTTCATC AGGAACAGAC TCCCGTACGA TTTGGATGA	17880
AAATGGAGCA GAAAAACTTC TTGGTCGAGG AGACATGCTC TTTAAACCGA TTGATGAAAA	17940
TCATCCAGTT CGTCTCCAAG GCTCCCTTAT CTGGATGAC GATGTTGAGC GCATTGTGAA	18000
CTTCATCAAG ACTCAGGCAG ATGCAGACTA CGATGAGAGT TTTGATCCAG GTGAGGTTTC	18060
TGAAAATGAA GGAGAATTTT CGGATGGAGA TGCTGGGGT GATCCGCTTT TTGAAGAACG	18120
TAAGTCTTTG GTTATCGAAA CACAGAAAGC CAGTGCCTCT ATGATTCAAGC GTCGTTTATC	18180
AGTTGGATTT AACCGTGCAG CCCGCTCAT GGAAGAACTG GAGATAGCAG GTGTCATCGG	18240
TCCAGCTGAA GGTACCAAACTCTGAAAGT GTTACAACAA TAAAAAAATA GCTTCTTCC	18300
AAGTTGGAG GGAAGCTATT TTAGTGGCTA TTGATTGCTT TTATTTCTG AAGTTGGCAG	18360
ATTTGGACTGT TTTTCGTTT CAGTAGCAGG TTTACTGAA GCAGGAGTAG AAGAGTCCTG	18420
AGTTGCTGTT TTCTGATCTT CTTTTCTC TTCTTGACG CTAGATTTG GTGTTTCTC	18480
TTGCTGTGTT TTTCTTGAC TAGTGTAGT CTCTTAGTT GGACTGGGTGT TTTCTTCTAGG	18540
GGATTCCTTT TGGATTTCTT TGACAAATGGT TGTCGCTGG CTTGCGTAG GTTCTTTTT	18600
AATATTTTG TTATTATCCA AGGCGTT	18627

(2) INFORMATION FOR SEQ ID NO: 114:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 2560 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 114:

TTAAATACGT TACCTTCCTT CTGCACGTT AGCAGGTAAG TCATTGAAAT TTAAAGATCA	60
AGATATTACA ATTGAAGAAA CGACTGAAAC AGCTTTGAA GGAGTTGATA TTGCTCTCTT	120
TTCAGCAGGT AGTTCTACAT CAGCTAAGTA TGACCACATAC GCAGTAAAAG CTGGCGTGGT	180
AGTAGTAGAT AATACATCTT ATTTCCGTCA AAATCCAGAT GTTCCCTTGG TTGTTCCAGA	240
GGTCAATGCT CATGCACTTG ATGCTCACAA CGGAATCATT GCCTGCCCTA ATTGTTCAAC	300
AATTCAAATG ATGGTGGCTC TTGAGCCGGT TCGCCAAAAA TGGGGCTTGG ACCGTATCAT	360
TGTTTCAACT TATCAAGCCG TTTCAGGTGC TGGTATGGGA GCAATTCTTG AGACACAACG	420
TGAACTTCGT GAAGTCTTGA ATGATGGTGT GAAACCACGT GATTGTCATG CGGAAATCTT	480
GCCTTCAGGT GGTGACAAGA AACATTATCC TATGCCCTTT AACGCTCTTC CACAAATTGA	540
TGTTTTCACT GATAATGATT ACACGTACGA AGAGATGAAG ATGACCAAGG AAACTAAGAA	600
AATTATGGAA GATGATAGCA TTGCAGTATC TGCAACATGT GTGCGTATTG CAGTCTTGTC	660
AGCTCACTCT GAGTCTGTTT ATATCGAAAC AAAAGAAGTG GCTCCAATCG AAGAAGTAAA	720
AGCAGCTATC GCAGCCTTCC CAGGTGCTGT TCTTGAAGAT GATGTAGCTC ATCAAATCTA	780
TCCCTCAAGCT ATCAATGCA TTGGTTCGCG TGATACCTTT GTGGTCTGTA TCCGTAAGA	840
CTTGGATGCA GAAAAAGGAA TTGACATGTG GGTGTTCTA GATAACCTTC TCAAAGTCC	900
TGCTTGGAAC TCAGTTCAGA TTGCTGAAAC TCTTCATGAA CGTGGATTGG TTCGTCCAAC	960
AGCCGAATTG AAATTGAAAT TAAAATAGTC ATATCGTTA GGAGTTCAAGA TGAACCTCTT	1020
CTTTGAAATA GAGAGGTGTT TTGGTGTCTT ATCAAGATTT AAAAATGT AAAATCATTAA	1080
CAGCCTTAT TACCCCTTC CATGAGGATG GTTCCATTAA CTTTGATGCT ATTCCAGCCT	1140
TGATTGAGCA TTTATTGGCC CATCATACGG ATGGAATTCT TCTCGCAGGA ACGACTGCTG	1200
AGAGTCCAAC TTTGACCCAC GATGAGGAGT TGGAGTTGTT TGCGGCTGTA CAAAAGGTTG	1260
TCAATGGACG CGTTCCCTTG ATTGCGGGTG TAGGTACTAA TGATACGCGT GACTCTATTG	1320
AGTTTGTCAA AGAAGTAGCG GAATTGGTG GTTTCGCAGC TGGGCTTGCT ATTGTTCTT	1380
ACTACAACAA ACCTCTCAA GAAGGGATGT ATCAGCACTT TAAGACTATT GCAGATGCTT	1440
CTGACCTTACCA ATTATTATC TATAACATTC CAGGGCGTGT AGTTGTCGA TTGACTCCAG	1500
AAACCATGCT TCGCTTGGCT GACCATCCAA ATATTATCGG TGTCAAAGAA TGTACTAGCT	1560
TGGCTAATAT GGCTTACTTG ATTGAGCACA AGCCTGAAGA GTTCTTGATT TATACAGGTG	1620
AGGATGGAGA TGCTTCCAT GCCATGAACC TTGGGGCGGA TGGGGTTATT TCTGTTGCCT	1680

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CTCATACAAA	TGGGGATGAA	ATGCACGAGA	TGTTTACTGC	GATTGCAGAA	AGCGATATGA	1740
AGAAAGCCGC	AGCAATTCA	CGTAAATTCA	TTCCTAAGGT	TAATGCTCTC	TTCTCTTATC	1800
CAAGTCCTGC	TCCAGTTAAC	GCAATTCTTA	ACTATATGGG	ATTGAAAGCT	GGACCCACTC	1860
GTCTACCTCT	TGTTCCAGCA	CCAGAAGAAG	ATGCCAACG	CATTATCAAG	GTTGTCGTAG	1920
ATGGCCACTA	CGAACGCAACT	AAGGCAACTG	TAACAGGGGT	CTTAAAGACCA	GATTACTAAT	1980
AAAGACAATA	AAATCCGGCT	CTTTGTCAAC	TGTAGTGGGT	TGAAGTCAGC	TAAGCTCGAG	2040
AAAGGACAAA	TTTTGTCCTT	TCTTTTTGA	TATTCAGAGC	GATAAAAATC	CGTTTTTGAG	2100
AGTTTTCAA	GTTCCGAAAA	CCAAGGCAT	TCCGCTTGAT	AAAGTTGATG	AGATTATTGG	2160
TCGCTTCCAA	TTTGGCGTTT	GAATAGGGTA	GTTGAAGGGT	GTTGACGATT	TTCTTTTGTT	2220
CCTTTAGAAA	GGTTTTAAAG	ACAGTCTGAA	AAATAGGATG	AACCTGCTTC	AGATTGTCC	2280
CAATGAGTCC	AAAAAATTC	TCCGGTTCC	TATTCTGAAA	GTGAAACAGC	AAGAGTTGAT	2340
AGAGCTGATA	GTGATGTTTC	AAGTTTTGTG	AATAGCTCAA	AAGCTTGTTT	AAAATCTCTT	2400
TATTGGTTAA	GTGCATACGA	AAAGTAGGAC	GATAAAAATCG	CTTATCACTC	AGTTTACGGC	2460
TATCCTGTTG	AATGAGTTTC	CAGTAGCGCT	TGATAGCCTT	GTATTGGGA	TTTCGATGA	2520
AACTGATTCA	TGATTTGGAC	ACCCACACGA	CTCATAGCAC			2560

(2) INFORMATION FOR SEQ ID NO: 115:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 11303 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 115:

TATTGGATTT	CCCTTGCAAT	CAGTTTATGG	GACAAGCACC	CCGCAGCGCA	GAGGAATCA	60
ACGCCTTCTG	TAGCCTACAT	TTTCAAACCA	CCTTCCCACG	TTTGCCAAG	ATTAAGGTCA	120
ACGGTAAGGA	ACCAAGACCT	CTCTATGTCT	GTTACAAGA	CCAGAAATCC	GGCCCACTAG	180
GAAAACGAGT	CGAATGGAAT	TTCGCTAAGT	TTCTCATCGG	TCGAGATGGG	CAAGTCTTTG	240
AACGCTTTTC	TTCAAAACAC	GACCCAAAAC	AAATTGAAGA	GGCGATACAA	ACTCTACTAT	300
AATTCAACAT	CTCACTATGA	TTAGGTTCC	TTAACCTGA	TGAATAGTGA	GATTTTTGAG	360
TGGGCTTTGA	CTTAAATAGA	AAAACACCCC	ATGATATGAA	ACATGAAGTG	TTGTAAAGTC	420
TATGTTGTAG	GTGCTTATTTC	CACAATTCA	ATGTGACCAAG	TGATAACGAA	TACCATACAG	480

832	
AATCTTCATA TACACTAAC AAATGACTTT CTAATTATTT CAATTAGTTT TGGCTAGTAA	540
ATATCATTTG CAACAAACGC CCTCTCAATT CCTTATCCTG ATGATGCAAG ATATTCATTA	600
AGTCATGAGA GTTTTTCGCA TTGATGAATT GATTTAACAA TCTATCTTTT AATTCAATG	660
GAAGAGAAGC TGTCTTTAGT AGTCTAAAAA CTTCGTCATT TAAAGATGTC CTTTTATTAT	720
CTTTCCATTC AAATTTAGCT GTATCATTCT TATTTGGCAA TTCATTATAA GACACATTG	780
TTCCCTTAAA ATGAATTCTA TGTTTTCTAT TGCTTGGAAAC GATACTAGAA TCTCCTTGTA	840
ATGCTAACTC TACCAATTCCC ATTTCCAAT CGATTGATAA TCTTGTGTTA TATCTTGAC	900
CATTTTGATC TTCAAGCATT TCAAAAGAAT GTTGTGTTCC TGGGAATACA TACCAATCTA	960
CAACTTCAGG TAAATCAACA CCCATACCTA TCTCAGAACC AACCAAGGGA ATGATTGCAC	1020
CACTTTTGCGC AAACACAGGC GTAGTCGAGA TGTCCCTATA AACACTTAAAC TTACACACCAC	1080
CTGTCGTATTT TTTCCTCTGAA AAGAAGTCAT ACCATTCACT TTCAGGGAAAC CATACTCTA	1140
CTTTTGCGAGA TTGGAATGTC AAATCCATCT TTTCCTACAAT GGAGGCCACC ATCAGTTCTG	1200
TTCCAAAAAA GTATTGGTTT GGAAACATTAT AGCTCTCATC ATTCTCTGGA TAGAAATAAT	1260
AGATTGGACT GATTAATGGG GCACCTTCCT CATGTGTCTG TACATTCTAGT GTATATAGAT	1320
AGGGAATCAT CTGATGTCTC AAACGAAGGT ATTTCTCTAT AATCTTAGAT GPTGTTCTG	1380
AAAAAAACCA AGGTTCTTTA CTATTAAG GACTTCTAGA ACTATGTAAT CGAGTAATCG	1440
GACTAAAAAC ACCAAACTGT AGCCATCTAG TTTGTAGCTC TTGTCATAAA TCCCCAACAA	1500
TATGTCCACC GATATCATGA CTCCACCAAC TATAACCGAT ATTAGATGCT GTGCTGTAA	1560
AATAGGGTTG AAATCTTAAG GAATTCCAAC TAATAATAGT ATCCCTGAA AAACCAACAG	1620
GGTAGCGGTG ACTACCAGGA CCTGCATATC TTGATAAAAT CAAACCACCT TCTGCATTTT	1680
TACAACATAC CTGATAGTGA TAATGGTTA AAAGCCAAAG TGGATCTAGC ATACCTTG	1740
TCCCTTGTG CCAGTCATC CACCAAAAT CTACTCCCTG CTTTCTAGT TCATAATGAA	1800
CATCTTAAA GTAGGCTTCC CTAAAAGAGG GATTAACAAAT ATCAAAATAA GCAGGTTCTT	1860
CTAGTTCTAC ATTTAACCCCC AACCGTTTTG CGATTGAGG ATAAGCTTCT TCATAAGCCC	1920
GTATCCCACAC AGCAGGATGG ACATTTAAGG AGAGTTTAGT CTTCTATCA TGAAGTTGTT	1980
GCAATAACTG TTCTGGATTT GGTATTAAGT TTCTATTCCA ACTATATCCT GTCCAGGCCAC	2040
TTCCAAAGCG AGCTGGAATG TCAGTTATAT GCCAATCCAT ATCTAACACA CCGATAGATA	2100
ATGGAATTTT CTCTGTTCA AATCTGTCTA TTAAATCCAA GTATTCACTC GACGTATAAG	2160
GCCAATATCT ACTCCACCAA TTGCCTAAAG CATACTTGG CAACAAGGGT GTTGAACCAG	2220
TCAAATGGTA AAAATCTCTG ATTGCTCCTC TATAATCATG CCCATAGGCA AAGAAATACA	2280

833

GGTCAATTG ATTTCTCTC TCAATATAAC CAGATTGTC ATCCCAAATA AATCCTTGAG	2340
AATCATCCAA TAAGGCTATA CCATTCGGC TAATAATTCC ATCTTCTAAC GAGATTGCTC	2400
CATCTGCCTT ATCCAGAGTC CGAGCTGTC CTTTAACGT TTCATAAGAT TCACCAAAAT	2460
ACCAGCGACT ACCATATACG GCAAAATTTC CTTTAATTC TATAAATAAA TTTTCGGCGT	2520
TAAATTCTCC TTATTAAGA TGCAGATGAA AATAGTCCGT CATAATATCT AGTACGTTG	2580
ATGTCTCGAT ATAATCTAAC GAAATTGCG CAAAATCTCT ATTATAGATA AGTTGTGTCG	2640
TTCTATCTC AAAACTTCCA GTTGAGAGT ATTCTAACCT TACTAGCTTG TCTGTTAATA	2700
CAGAGATTG ATAAAATCT CCTTAAAAA TTTCATTTT GTTTCCCTCC TTTTATGGTA	2760
GCATAAAAAC AGAACGCACC ATTTTGATG CGTTTTCAT TATTCTGAAT GCAATGTTCT	2820
ATCTGTTATA TCTATGACAA ATAATAGTC ATTGAAAAAA TGCAGTGGAC AAAATATCTT	2880
TTAACAAACC AAGAGTTTAAAGAGTTA TCACTTTCA ACTTTCTAA GCTTATGCGAG	2940
TTGTGAAACA AACTACTTTT AAACATTAA CTAAGATAGG ATTGATAAT AATTCAAAC	3000
TCTTACTAGC AATCATAACG TATTCAAGCT CACGTGCTTT TTTCCCTCCT GCTTATTCT	3060
TAGAACTGAA GAACCCGGAT CGGTATATAA ATTATCCGGA TCAACATAGT CATAAGATT	3120
ATAACAGTTG CGCTTCATTA AGTCATCCCC AGAGCAAGAG CTTCATCTCG TAATTTTCA	3180
ACATCACTAA CGTAGGTCG CCATCCTCA ATCATATTG TACTTAAAC ATACAAACAA	3240
CTCTTAAAAA CGGATCGGTT TTCAAAAGCT ATTCCCCTGA TTGTCATCTT TTCTTTATCT	3300
ATATCTAAGG ACATATGCTA CCTCCTTTAG ATACATTATA CCATGTTCT CTGTAGCTTT	3360
TAAAAATTAAATTTT ATTTGTTG TCATATCTAA GTTTTCAGCA CGCTTATCCT ATTTTATAAG	3420
CCTCAAACCC AAATATAAA CGCATTCTTT TTGCTTTTTT ACTATTGTAT CGTATTCTAC	3480
GATAACATAC TTACTTTAT TGTCTTTTA AATAACAGCA GTTCCCTGTT TATCAACTAT	3540
TCGAACACT TTCTATTTG CTTCATACCC TACATAGCGA AAAATATGA AAAAGCAGAG	3600
AAGAAATATCT TAAAAAGACC TCTTCACTGC TAATATTAAC ACTCATTATT TAAACTATAT	3660
GGATTCTATC ATCGAGTATA CTTTTTACT TATTAGATACT CTGCTCTC TTTCACCAAT	3720
TTTGATCAT ATACACGGAT GAATGGAAGA TAGACTAGGA ATGCTGCAA TGACACACT	3780
AGAGCAACTA ATACAGCTCG AAGATCTGCT GTCCCTAAGA AAGCTCCAAT CCCTACTGGA	3840
GTTGGCCATG GAACCTGTGC GATAATTGGC TTAATAAAAGT TTGAGAATT CGCTACGTAA	3900
TAAATAGTAG CAGTAACCCT TGGTGCTAAA ATAATGGTA TAGCCAAGGC TGGATTATAG	3960
ATAATAGTA ATCCAAAAT TAATGGTTCA TTAATATTAA ATAAGGCTGG AACTACAGAT	4020

834	
GCTCGTCCTA TTGCTTAAG CTGTCAGAT TTAGAGGCAA AAGCAATATA TAAACATAGT	4080
CCTAAAGTTG CACCGAAC ACCTGCAATT ACAACATAT TAGAAAATTC ACCTGCAACA	4140
GCGAAGTGC CGCCAGCAGC ATTTTCAGCC ATGTTAGCAA GAGCAATTGG ACTAACAAAT	4200
GCAAAAACAA TGTTCGCACC GTGGATACCT ACAATCCAAA GTAGTTGAGT CAATAGATAA	4260
ATAATCATTA AACCAATCCA CGAATTAGTC AGATTGGATA CAAAACAAA TGGAATTGCA	4320
ATGACTTTAA AAATATCTGT TCCCATTGCT ACAAGAAGAC CGTTGATAAA GATAACAACA	4380
AATGCAACAA CAAATCCCGG AACCAAAGCG GTAAATCCAC GAGAAACTCC TTCTGGAACA	4440
GCTTCAGGCA TTTTAATAAC CCAATTATGT TAAACACACA TACGATAAAAT AAGAACAGTC	4500
ACAATTGCCA TAATGATTGC GGTAaaaATC CCTGTTGTCC CAAAACGTGC GACTACATT	4560
CCCATTGCCC ATCCATCTGC AATTACTGCA CCTTCTTTTA GACTGTGTCAC AGTCTTCATC	4620
ATTCACCAT CAAAATGAT TTGCGGTACT GTCATGACAA AAGCCATCAA GGCAAGCAAG	4680
GCACCAATTAA GAGGATTCAAT TTGAGTTCT TCTTCCTCTG CATAAAATTTT TGTCAATTCA	4740
TATGCAAGTG ATAGAACGAA ATAAAGAGAT AGAGAACCCA TAGTCGCATA GTTTGCAACC	4800
ATGTAAAGTG ATGTGAATT ATCAAATGAA GCAGAGAAAA TATCTGCCAC AATTGGCCAA	4860
AATGAGAAAG CTTGTGGCAA AATACTGAAT ACCAAAAACA TTGATCCTAC AATAGTAAAT	4920
GGTACAGCAG CCATACCTGC AGCCGTGATA GCACGTACTA CTTTAAACTG AGCAAGTTG	4980
CCCATTGGTC CCATAACATG GTTTCAAGA AAACCAAACA ACCCGTTTG TTGATCCATA	5040
AATAGACCTC CTTAATTTAA CATAATAATT TTACTTTCT AAGACTAGT TTCAAATACA	5100
AATTATACTA GATCAGGATT ATAAACTAAG TGAGTTCTTT TCCAATTGGA CAAATTGTTG	5160
ATAAGCCTTA TCTGTTGCTT TATAAAATTTT TTAAATTCTT CTAATGTCTA ACAAACTCAG	5220
AACTAAACCT AATAGAACGAA CTACAAAAAC AAATAAACGT GCTACTTGGT TATTTCAAA	5280
AATCGGAAAA AGATTCTAA ACCAACTTGT CCAAGTTAA ACAAGTAATC CTATTGAAAT	5340
AAGCATTTGT ATTCTAACAA ACATTAGTGT TATTCCCAAC TTTCTTTCC TATTTCCATA	5400
AAGTTAAAT TGTTCAACAG TTGCTAAAAT AGAAAATACT ATGAGCATAA TGGGGAAAAT	5460
AATAATAGGC GAGGGACTAA TAAACTGACT CAAAAGCCAA TAAATATTCC CAAAAAAGAA	5520
GAGTGTCTATT GAATAACGTA GAAGAACGATA TCGATTGAAA AAAGTATTAG TTAGAGCCAT	5580
CTCTCGACGT TGTTGTCCTA TCTTTGTCG TTCTTTTTA TCCATATCAT TTCTCCTTA	5640
TATAACAAACA CATATTTAGT TAACTTTCTT ATAAAGAGCT AACATTTCCCT TTGCTACTTC	5700
TAATAATGTC ATAGTGGTCA TTAAATGATC TTGAGCATGT ACCATGATAA TTTCAATT	5760
AATTTCACCT CCACCTTGCGT ATTCTTGCAA GAGTTGGTT TGTGCATGAT GCGCTTCAG	5820

835

AATTATCTCA	TTTGATTGAT	TTAATTTACT	TTCTGCATCA	TCAAAACTAC	CTTCTCTCAT	5880
TTTTGCAAAT	GCTTCATGTA	TTTCTGACCT	TGCATTTCCC	GAATGCAGGA	TAATTTCAAA	5940
TGCTGCAACC	TGCAGTTCCCT	CTTGATTTCAT	ATAAACCTCC	TATTTTATCT	TCTCAAATAT	6000
GTAAATAAAA	TCTTCAAAGT	TATTGCAAGA	TATTAGCTGA	TTTTGCAATT	CATCATTCTC	6060
TGTCAGAGAG	ACTATCTTTT	TAGTCACAGT	TGCCAAACCT	TCGTTCCCAT	ATATTGATGG	6120
AGATAGAAGA	AATACTAGCT	GGACATGTGA	ACTTTGATTA	TCCCAGAGTA	ACGAATCTTT	6180
ACAAATTGCA	ACCGAACCT	TTCCCTCTGT	ACCAAAGGC	TGAATAGGAT	GCGGAACCTGC	6240
AATTTTTCA	GAAAAAACAA	CTGAACCTAA	TTCTTCGCGC	TGTTTAATTC	CATAAAGTAA	6300
AGATTGTTCA	AACTCATTG	ATTCACCAAC	AGATAAACTC	TCAACCATCT	TTTCAAGTAA	6360
ATTTACCTTG	TCTGATTCA	TACATATTAA	AAAGTTTCT	TTACTAAAAT	ACTGTCTAAA	6420
GCCGTTGTTT	TCAAATTGTT	TAATCTTTGA	TGATTGTACA	TAACTAGAAA	CTTGCATCTA	6480
ATCCATAGCT	TTTCTAATCA	TTTCCATCTC	ATCACTCTTA	AGAAACACAC	TAACTTTAAA	6540
AACTGGGATT	TGAAATATA	GATTGATAA	ATCAATAGCT	GACACTATAA	AATCTATTCC	6600
TTTAAGTTTT	TCTTGATTCA	ATTCA TAGTA	GCCTATTACA	TCAACAACTT	CTACTCGCTT	6660
CCCAAACCTCC	GTTTCCAAAC	GATTCTTAA	CATTTGGGCT	GCACCAAATC	CTGTTGCACA	6720
AATACCAAGA	ATATTAAC	TAGTACTCTC	TTTGCTACGT	TCCATAGCAG	CTAAAAGTG	6780
AAGACTTACA	TATGCTACTT	CATCATCTGA	TATTGTCCAC	TCCAAGAACT	TGTCCATATT	6840
TGCAAGAATT	TCTCTAGTCA	AAAAGAATAT	ATCACTATAA	TTCTGTTAA	TTTCATCTAC	6900
CAAAGGGTTA	TTTAAGGTA	TCCGGTTTC	AAACGTACT	TGTAGTGTCA	TTAGATGAGT	6960
TATCAATTCT	TCAATTAGTT	GGAAATCTGA	AGAAAAGTTA	TACATATCAT	CTAATCCTAA	7020
ATTCTGAAAT	GTTTAAATA	AAAGATTTTT	AAAACCTCT	TCAGAAATAT	TCTTCTGATT	7080
TTTTGACAT	TGTTGACTCT	AGCTAACAA	ATGCAAAGTA	ATGTAGTCTA	TTTCCTGAAC	7140
TGGAAATTCC	TGATTGTTA	CTTCTCTTAC	TTTAGAAAGA	ATTCTTTGGG	CAACCTTTCT	7200
CTCTATGCA	TCATCAGTCA	TCTGACAGTC	TATATTTTT	ATTCAAATC	CGGATTTAA	7260
ACGAATCACA	GACAATGCTA	TGTGAAC	AAATTCTGT	AGTACAAAAT	CAGATAGTT	7320
TAGGTTGGCC	TCTTGGCATT	CATCCAAAC	AAATTCTAGCA	AAATTCTTCTA	ATGGAACAGT	7380
TTGATCAAAA	AAAGTAAATT	TTACATAGCA	ATGTATTGTT	TTAAAAAAATT	GATTCTCTAG	7440
GAAATAATT	ATGATAAAAC	GTCGTTTATC	ACGTTCCCTCG	CCTGAGACAT	AAACTCCTTT	7500
ATTCGCCCTA	CTCTCAATGG	ACAAATTATA	CTCTGATAAC	ATCACTCGTA	TCTTCTGAA	7560

836	
ATCATGAGAT AATGTTGAAC GACTAACGTA AAGTTCATCA GCTAAATCAT CAAAAAGAAC	7620
TGGAACCTGC TCAAAATAATA ATTTATTTAA GATAAATACT AAACGATCAT CACCTTTGA	7680
AACCGCAGTT TTCTGTATAGT CTTCTTCAG TTCATAAGTT TGTCTAAACT CCTGGTAAGC	7740
GCCTTGATTG TCAAAAAATA TTTGATAACCC TTGACCTTGT TTTGAAATCA ACCGGACTCC	7800
TTGAATAATC ATTGTCTTCT CAATTAATTT CAGTACATTA CGGACAGTTC TATCTGAACA	7860
GGATAAAATAT TCTGCCAGTT CTTTGCCTGT AACAAAACGT TCCTTATTTT TTATTAAAAAA	7920
TTGAAGGATA TCTTTCTCTT TAATGTTAA CACATTCAATT CCCTCCTAAA ACGTATGTTT	7980
TCATATATTG AAGCATATTA TACACTTAA TCAGTTATA TCAAACCTCAA AACAAATTTAT	8040
CTTAACCTAA ATATTATTTG ACATTTCATG TGTTCATCAA ATATTCTCAA GAATCAAATT	8100
AGCCATTTT TCAATTCCC TTGGAATAGG AATATAGGCT TGAGGAGGTA TTTGTACAAC	8160
TGGTTTCTCT GCTTTAGAAC CAGCCTCTTC AAATIGCTTA AAGTACATTT TTGTTTGAGG	8220
ACTGACAAGA TACAAATCAA AAGCTGCTGC TGCGATAGCT TTCCCTCCTT CAGTAGCACT	8280
AATAGCATCA ACTACAATAT CTTTCCCTTT TCCTTTAGA AACTCTGTTG TTTTCTGTGC	8340
CATAAGTGAT GAAGACATTC CTGCTGCACA AATAATTAAA GCTTTGCCA TAATATTTTC	8400
TCCTTTCTT AAATCCAATC AAAGCTGTGC TAAGTTGGCT TATTTGTTAT CTATTTTTAT	8460
TATAAAATAA AGCGTTTCCA ATGACAATTC CCTCATTTTC CTAAATGATA TGGAAAAAAA	8520
TTATTTATAC TTCAATTAT AAAATAAAAT TATTCCTGAG AGTAGAAATG AAACACTATT	8580
TGCTAAATC AAAGGCAAGT CTCTTACG AATACCATGA GCAAGCCACA ATGCAATACC	8640
AATAACTTGC ATAACATACA TACCTAGAGC AATAGATCCT GTGCTCTTG TCTTAACTAC	8700
ACGAAAAACT TGTGGAAAA ATGCAAATGT TGTAAAATT GCTGCAATAC TTCCAATCAT	8760
ATGTCACCTC AATATGCTAA ACAAACTGAG AATAATCTCA GTTTGTTTAT ACTATTCTAC	8820
TGATTCAACCG TTAGATGAA TAACCTCCTT ATACCAGCCA AAAGATTTT TCGGGGAACG	8880
ATTATAACTT CCCTTCCAT TATCATCTTT ATCTACATAA ATAAAGCCAT AACGTTTCCG	8940
CATTTCACCG GTACCAGCTG AAACCAAATC AATACATCCC CATGGAGTAT AACCCATTAA	9000
ATCAACACCA TCTTCAACTA CAGCCTTTT CATTTCACGA ATATGGGCAC CTAGATATT	9060
AATTCTATAA TCATCATGTA CCATACCATC TGCTGCAACT TGATCTATAG CTCCAAAACC	9120
ATTTTCACCA ATAAAGAGTG GTAAGTGATA GTGGTCTGTA AACCAATTAA ACGCATAACG	9180
CAAACCTCT GGATCAATTG GCCACTCCC TTCAGAAGCC TTAACATAAT TATTTTCAC	9240
TAAATCTTCT GTTCAAGAT AATCAAAATA AGGATTATTT TCACGATGAG AGTCGATAGC	9300
AAAGGACATA TAGTAACTGA AACCAATGTA ATCTACAGTC CCACCAAGTA AATCTTCTTT	9360

837

ATCCTGGCA GTAAAATCAA CTGAAATACC TTTCGTTCC CAATACTTGA AAATATGCTC	9420
AGGATATTTA CCTAAAACAT GCACATCAGC AAAATAATAA CGCTTCTGCA TAGCTTCAT	9480
TGCCATTAAG ATATCCTTAG GATTGCAAGT AACTGGATAA ATTGGACACA TCGCAATCAT	9540
ACAAACCTATT TGAAAATCTG GATTAATCTC ATGACCAATT TTTACAGCTC GTGCAGAAC	9600
AACTAATTG TAATGTGCTG CTTGATACAT AATTGCTTCT CTATTATCAC CTTCCCTCATA	9660
TACAATACCT GAGTTAGTAA ATGGTGAAA ATCTTCCTGA TAATTCGCTT GATTATTGAT	9720
TTCATTGAAA GTCATCCAAT ATTTAACCTT ATCTTTGTA CGTTTAAATA CGACTTCTGC	9780
AAAACGAGCA AAGAAATCAA TCAATTCCCT ATTTTCCAA CCACCATATT CGGTCACTAA	9840
GTGATAAGGC ATTTCAAAAT GAGATAGAGT GATGACAGGT TCAATACCAT TCTTTAAGCA	9900
TTCATCAAAA AGATTATCAT AAAACTGTAA TCCTTCTTCA TTGGCTCTA ACTCATCACC	9960
TTTGGAAAG ATACGTGTCC ATGCAATAGA GGTACGGAAG CACTGAAATC CCATTTCAAGC	10020
AAAAAGTGCT ATATCTTCTT TATAACGGTG ATAAAAATCT ATCGCCTCAT GATTGGATA	10080
ATATTTACCC TCTAAAACTC CCAAAGTAAT TTCACGAGCT ACTCCATGAC GACCAGCAGT	10140
CATAACATCA GCAACACTAA TTCCCTTGCC ACCTTCTTGC CATCCACCTT CAAGTTGATG	10200
AGCAGCAACA GCACCACCCCC ATAAAAATCC ATCTTTAAA GTAGTCATCT TTTTCTCCTC	10260
TGACTTTGAT ACTCTTATTA TAAACCTTAA ACCAAAAGAT GAAAACGAT TCTTTTCCT	10320
TATGTTAAG GAAAGAAGTA ATTTTTAATG GAAATAGAAC AATATCTTCT TGTATTCTCG	10380
TAATGATATC TTACGATTT TCAATACCTT CAAACTACAA AAACCTCTCAC AATAATTCTA	10440
ATCCCTGTG TCTATAAACG ACTTATCGCT TTCTGGCATC CCAGAACATC CTTCTATATA	10500
ACGTTCAACT TGCATCTGCA AGTGATATT TTTCTTAAA TCTAAGATTT TCTGCATTGT	10560
CTTGATTGA TAATGTTTAT CTAAAGTTTC TTGATTATC CACTGATCAA TAAGGAGAAT	10620
AGTTCCCTCT TTTCAATTG GTAAAAAATA TTGATTTTC AAGTTACCTT TTTGATPTCT	10680
AATTTCTTAA ACAAGGCCAC TATCAAGCAT TTCTCTTGC AACTTTATG CACTATCTCC	10740
ATCACCTTTA TAATATACAT GAATAGTCAA TGTCACTTAA TATCCTCCAA AATCATCCTT	10800
CAATTTAAA AAAACAAGTT TAGATGAGGA TCTAAACTTG TTTTTATGA ACTAATTATC	10860
TAACGTTTCG CCATTACTTT CAATCACTTC TTTATACCAA TAAAATGATT TTTCTTATA	10920
GCGATTATA GTCAATTGAA ACAAGAGCAG GACAAAAGAG CCTCATAAAA GGTATTGCAA	10980
CTTGGTAATA CCTTTTGAG GTGCTTTTG ATATGAGCCC ATGTTTCTC AATAGGATTG	11040
TACTCAGGTG AGTAGGGAGG AAGAGGTAA AGTTTATACC CAAACTCTC ACACAAGAGT	11100

838	
TCTAGCTTCC CCATTCTATG GAATCTTGCA TTATCCATAA TAATAACCGA TGGTGTGGTT	11160
AATGTTGGTA AGAGAAACTT CTGAAACCAA GCTTCAAAAA AGTCGCTCGT CATCGTCTCT	11220
TCGTAAGTCA TTGGAGCGAT TAACTCACCA TTTGTTAGAC CTGCAACCAA AGAAATCCTC	11280
TGATATCTTC TTCCAGATAC TTT	11303

(2) INFORMATION FOR SEQ ID NO: 116:

- (i) SEQUENCE CHARACTERISTICS:
- (A) LENGTH: 3112 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 116:

CCTTAGATTG CCACTTGCCA GAGGAATTGA TTGCCCAAC GCCCCTTGAA AAACGTGATG	60
CCTCCAAACT CCTCATCGTC AACCGTGAGA CAGGAGAAAT GCAAGATAAA CATTTCAC	120
CTATTATTGA TATGCTGGAA CCTGGTGATG CCCTTGTAT GAACGACACC CGAGTTCTCC	180
CTGCCCGCCT CTATGGTCAA AAAGTGGAGA CAGGAGGTCA TGTGGAACCTT CTCCCTCTTA	240
AGAACACTAG TGGAGACGAG TGGGAAGTTC TGGCTAAACC TGCCAAACGC CTCAAGGTCG	300
GTACTCGTAT CAGCTTTGGT GATGGCCGCC TCAGCGCTGT CGTTACAGAA GAATTGACCC	360
ACGGGGGACG CATTGTCGC TTTGAATACC AAGGAATTTC CCTAGAAGTC TTGGAAAGTC	420
TGGGAGAAAT GCCTCTGCCA CCTTATATCC ACGAAAATT AGATGACCGT GAACGTTATC	480
AAACCGTCTA CGCCAAGGAA AGTGGCTCTG CTGCAGCACC GACTGCTGGT CTTCACTTCA	540
CCAAAGAACT GCTGGCAGAA ATCCAAGCTA AGGGTGTCA TCTAGTCTAT CTGACTCTCC	600
ATGTCGGACT CGGAACCTTT AGACCTGTTT CTGTGGATAA TCTGGACGAA CACGAAATGC	660
ACTCAGAGTT CTATCAACTT TCTGAGGAAG CTGCTGCCAC CCTTCGCTCT GTCAAAAAAA	720
ATGGTGGTCG TGTCTCGCT GTCGGAACCA CTTCTATCCG CACCTTGGAA ACTATTGGTT	780
CCAAGTTTGA TGGGCAAATC CAAGCAGATT CTGGTTGGAC CAATATCTT ATCAAACCTG	840
GGTATGAGTG GAAGGTCGTG GATGCCCTCT CAACCAACTT CCACCTGCCA AAATCAACTC	900
TGGTCATGTT GGTTCTGCC TTTGCAGGCC GTGAATTAGT CTTAGATGCC TACCACCTT	960
CCATCCAAGA ACACCTACCGC TTCTTCAGTT TTGGTGACGC CATGTTTATT TATTGAGAAA	1020
GAATTCTCT AAATCTTCTA ATACCAATAA ATCGCTAAGA TATTATTCGA AAGAACATCT	1080
ACAATTGAAA CTCTAGCTAG CTGTAGAAGA GGCCTAGTAC ATTGAAATTAA AAATGCTTCC	1140
CCCTAGCTTC GAAAATATTG CCATAGATTG CGTTGACTCT CCAAATTGAT TCATCTATAT	1200

839

TTTATTCAG CTTCCCTAC TTTCTCGCT GTTTGTAAT CAAAATGCAA GACACATGAG	1260
TAGCACCATA TTGGTTACTC TTATCTGTCC TCTCAAGAGA CTATTATGAG TTATTCAGA	1320
ATCATTCACT ACTTTGACCC TGACTCTCCT TAGTCTAAA ATCAAAGACT TATACTCTTC	1380
AAAAATCTCT TCAAACCGCG TCAACGTAC CTTGGATTAT ATATGTGatC TGaCTTCGTC	1440
AGTTCTATCT ACAACCTCAA AGCAGTACTT TGAGCAACCT GCGACTAGTT TTCTAGTTG	1500
CTCTTGATT TTCATTGAGT ATTAAACAAA AAGTGAACAA ATCTGAATTC TAATGTACAG	1560
AAGACTAGGC TTGTTCACTT TTTTATAGTC GCTATAAGAT GACCTTATCT ATAGCTTTT	1620
ATATATAATT ATATATTCAAG ACATACTATT ATCAATTGG TCGCAGGGAG GAATCTGTTA	1680
ACGCACCCAT TCACCAATT CATTGACTCT ATAGCCATCT ATACTTGTAT TGACCGCTAA	1740
CTCACCCGAT GTATTTACAT AATACCATT ACCACCAACT TGGAACCATT GATTGACTTT	1800
CATAGAACCG TTGCTGTTGA GGTAGTACCA TGAACATTAA ACTTGTACCC AACCTGTTGC	1860
CATGGAACCA TCAGTATTAT AAAAATACCA CATAACCATT TCTTGTTCAG AGTCTGTTGT	1920
TGGAGCAACT GCTTTAGCTG GTTCTACTGC TACATCTGTT CCTTGGTTAG ATGTAACAGA	1980
TACAGGATAC GAAGGAATAG ATGATTGCTC AGGAACAACA ACTTTTTCAG GTTCTCTCGT	2040
CCCTCTCCTT ATACGTCTT TTACCATCTC TTTAGTAATT TGACGAGAAG TAGTTCTTC	2100
AATTGTTCCA TCACGTTCAT CTACAGTATA GATTGTAGTA AGAGTAATT ACCAATTCT	2160
CCTACTTCTT CTACTTCTG ACTTTTATCA AGAGTTGGC CATCGAGATA TTCTGTTCG	2220
ATTGGAATT CTTGGACAAG AACTTGGGGC TTGGTTCTTT TTTAACAAAC TCTTGTGTTGA	2280
GAGTCTTTT TTGACTTAA AGTACTCTCA GTTACTTGTG CACTCTTCC ATCTACATTA	2340
TAAGTTATCG TTGTAACTGT TTTCCCATTC TTTCCTAGAG TAATCTCTG CTCTGTCCT	2400
GCAGAAAGGT CATTGCTGC TTCATATTAA GTAGCAAATG GAACAAGAAC TTCTTCAACC	2460
TTGCTTTAG CTGGAACCTT GATAACTGTA TCCGTGGCTT CTTTTCTATC AACAGTAACC	2520
TGTTCGGTA CATAACCAGT CTCTGGATTA ACATCGTAGG TCCTTGTGCGT AGTTACATAG	2580
CCATCCTCTC CATCAATTGT AACAGGATTT TCACTACGGT CTTTTGTTTC ATCTTTTCA	2640
TAACGAATTG CGGTACTTGA AATTTTCTTG GTTACTACCT TAGGTTTTAGT CGCTACTTTT	2700
ACAATAATAT CCCCATGGTC AGCGTCATCA TACTCTATT CCTCTTCTTT ATCTCTAGTA	2760
TCATCTCTGA CATATTGAAT CCCATCAGCA GCATGAACAA AACTTGTATT CAGATTCCTC	2820
CTAAAAATAA AGTTAGCCCG ATTACCGCAG AACCAAAAT CTTTCCGAGT TTACGTATTG	2880
CATAGCGCTT ATTAGTATTA GATTGGCCA TTACATCCTA CTTCTAGTAT ACCATCTTTT	2940

CTATCAAACG TTAAACAATA TACGTTATAT ATAAAATAGA CTTAGAATGA TATATTGATT	840 3000
ATTGAACTAA CACTTTAACT ATATCGTAAT CAATCTCATA TATAAAGGAT TGCGACACATC	3060
TTATCTAAAT ACATGCGAAT ATATTTAGAT ACAAACATTC CAACTTGATA AT	3112

(2) INFORMATION FOR SEQ ID NO: 117:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 4327 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 117:

CCCAAAATC TCCTCAAACC ACGTCAGCTT CGCCTTGCCG TAGTATGGTT ACTGACTTCG	60
TCAGTTCTAT CCACAAACCTC AAAACAGTGT TTTGAGCATC ATGCgGCTAG CTTCTTAGTT	120
TGCTCTTTGA TTTTCATTGA GTATAAAAAC AGATGAGTTT CTGTTTTCTT TTTATGGACT	180
ATAAATGTTCA AGCTGAAACT ACTTTCAAGG ACATTATTAT ATAAAAGAAT TTTTGAAAC	240
TAAAATCTAC TATATTACAC TATATTGAAA GCGTTTTAAA AATGAGGTAT AAAAAATTAA	300
CTAACCGCTTA TAAAAAGTGA TAGAATCTAT TTTTATGTAT ATTTAAAGAT AGATTGCTGT	360
AAAAATAGTA CTAGCTATGC GAAATAACAG ATAGAGAGAA GGGATTGAAG CTTAGAAAAG	420
GGGAATAATA TGATATTAA GGCATTCAAG ACAAAAAAGC AGAGAAAAG ACAAGTTGAA	480
CTACTTTGA CAGTTTTTT CGACAGTTT CTGATTGATT TATTCTTCA CTTATTTGGG	540
ATTGCCCCCT TTAAGCTGGA TAAGATTCTG ATTGTGAGCT TGATTATATT TCCCATTATT	600
TCTACAAAGTA TTTATGCTTA TGAAAAGCTA TTTGAAAAG TGTCGATAA GGATTGAGCA	660
GGAAGTATGG TGTAATAGC ATAGGCTGAT GTCCATCATT TGCTTATAAA GAGATATTTT	720
AGTTAATTG CAGCGGTGTC CTGGTAGATA AACTAGATTG GCAGGAGTCT GATTGGAGAA	780
AGGAGAGGGG AAAATTGGCA CCAATTGAG ATAGTTGTT TAGTCATTT TTGTCATTTA	840
AATGAACTGT AGTAAAAGAA ACTTAATAAA AGACAAACTA ACTGCATTTT CTGGAGTAAA	900
TGTCTTATTT CAGAAATCGG GATATAGATA TAGAGAGGAT CAGTATGAAT CGGAGTGTTC	960
AAGAACGTA GTGTCGTTAT AGCATTAGGA AACTATCGGT AGGAGCGGTT TCTATGATTG	1020
TAGGAGCAGT GGTATTTGGA ACCTCTCCTG TTTAGCTCA AGAAGGGCA AGTGAGCAC	1080
CTCTGGCAAA TGAAACTCAA CTTTCGGGGG AGAGCTCAAC CCTAACTGAT ACAGAAAAGA	1140
GCCAGCCTTC TTCAGAGACT GAACCTTCTG GCAATAAGCA AGAACAAAGAA AGGAAAGATA	1200
AGCAAGAAGA AAAATTCCA AGAGATTACT ATGCACGAGA TTTGGAAAAT GTCGAAACAG	1260

841

TGATAGAAAA AGAAGATGTT GAAACCAATG CTTCAAATGG TCAGAGAGTT GATTTATCAA	1320
GTGAACTAGA TAAACTAAAG AAACTTGAAA ACGAACAGT TCACATGGAG TTTAAGCCAG	1380
ATGCCAAGGC CCCAGCATTG TATAATCTCT TTCTCTGTGTC AAGTGCTACT AAAAAAGATG	1440
AGTACTTCAC TATGGCAGTT TACAATAATA CTGCTACTCT AGAGGGCGT GGTCGGATG	1500
GGAAACAGTT TTACAATAAT TACAACGATG CACCCTTAAA AGTTAACCA GGTCAGTGGA	1560
ATTCTGTGAC TTTCACAGTT GAAAAACCGA CAGCAGAACT ACCTAAAGGC CGAGTGCGCC	1620
TCTACGTAAA CGGGGTATTG TCTCGAACAA GTCTGAGATC TGCGAATTTC ATTAAAGATA	1680
TGCCAGATGT AACCGATGTG CAAATCGGAG CAACCAAGCG TGCCAACAAT ACGGTTTGGG	1740
GGTCAAATCT ACAGATTCGG AATCTCACTG TGATAATCG TGCTTTAACCA CCAGAAGAGG	1800
TACAAAAACG TAGTCAACTT TTTAAACGCT CAGATTTAGA AAAAAAACTA CCTGAAGGAG	1860
CGGCTTTAAC AGAGAAAACG GACATATTG AAAGCGGGCG TAACGTAAC CCAAATAAG	1920
ATGGAATCAA GAGTTATCGT ATTCAGCAC TTCTCAAGAC AGATAAAAGGA ACTTTGATCG	1980
CAGGTGCAGA TGAACGCGT CTCCATTGCA GTGACTGGGG TGATATCGGT ATGGTCATCA	2040
GACGTAGTGA AGATAATGGT AAAACTTGGG GTGACCGAGT AACCATTACC AACTTACGTG	2100
ACAATCCAAA AGCTTCTGAC CCATCGATCG GTTCACCAAGT GAATATCGAT ATGGTGTGG	2160
TTCAAGATCC TGAAACCAA CGAATCTTTT CTATCTATGA CATGTTCCCA GAAGGGAAGG	2220
CAATCTTTGG AATGTCTTCA CAAAAAGAAG AAGCCTACAA AAAATCGAT GGAAAAACCT	2280
ATCAAATCCT CTACCGTGAA GGAGAAAAGG GAGCTTATAC CATTGAGAA AATGGTACTG	2340
TCTATACACC AGATGGTAAG GCGACAGACT ATCGCGTTGT TGAGATCCT GTTAAACCAAG	2400
CCTATAGCGA CAAGGGTGAT CTATACAAGG GTGACCAATT ACTAGGAAAT ATCTACTTCA	2460
CAACAAACAA AACTTCTCCA TTTAGAATTG CCAAGGATAG CTATCTATGG ATGTCCTACA	2520
GTGATGACGA CGGGAAAGACA TGTCAGCTC CTCAAGATAT TACTCCGATG GTCAAAGCCG	2580
ATTGGATGAA ATTCTTGGGT TAGGTCCTG GAACAGGAAT TGACTTCGG AATGGGCCTC	2640
ACAAGGGACG GATTTTGATA CCGGTTTATA CGACTAATAA TGTATCTCAC TTAGATGGCT	2700
CGCAATCTTC TCGTGTCACTC TATTCAAGATG ATCATGGAAA AACTTGGCAT GCTGGAGAAG	2760
CGGTCAACGA TAACCGTCAG GTAGACGGTC AAAAGATCCA CTCTTCTACG ATGAACAATA	2820
GACGTGCGCA AAATACAGAA TCAACGGTGG TACAACATAA CAATGGAGAT GTTAAACTCT	2880
TTATGGCTGG TTTGACTGGA GATCTTCAGG TTGCTACAAG TAAAGACGGA GGAGTGACTT	2940
GGGAGAAGGA TATCAAACGTT TATCCACAGG TTAAGATGTT CTATGTTCAA ATGTCTGCTA	3000

842	
TCCATACGAT GCACGAAGGA AAAGAATACA TCATCCTCAG TAATGCAGGT GGACCGAAC	3060
GTGAAAATGG GATGGTCCAC TTGGCACGTG TCGAAGAAAA TGGTGAGTTG ACTTGGCTCA	3120
AACACAATCC AATTCAAAAA GGAGAGTTG CCTATAATTG GCTCCAAGAA TTAGGAAATG	3180
GGGAGTATGG CATCTTGTAT GAACATACTG AAAAAGGACA AAATGCCTAT ACCCTATCAT	3240
TTAGAAAATT TAATTGGGAA TTTTGAGCA AAAATCTGAT TTCTCCTACC GAAGCGAACT	3300
AGAGAGATGG GCAAAGGAGA GATGGCAA GGAGTTATTG GCTTGGAGTT CGACTCAGAA	3360
GTATTGGTCA ACAAGGCTCC AACCCCTCAA TTGGCAAATG GTAAAACAGC GACTTTCTA	3420
ACCCAGTATG ATACCAAGAC CTTGTTGTTT GCAGTAGATA AGGAAGATAT CGGACAGGAA	3480
ATTATTGGTA TAGCTAAAGG AAGCATCGAA AGTATGCATA ATCTTCCTGT AAATCTAGCA	3540
GGTGCCAGAG TTCTGGCGG AGTAAATGGT AGCAAAGCAG CGGTGCATGA AGTTCCAGAA	3600
TTTACAGGGG GAGTTAATGG TACAGAGCCA GCTGTTCATG AAATCCAGA GTATAAGGGA	3660
TCTGATTGCG TTGTAACCTCT TACTACAAAA AAAGATTATA CTTACAAAGC TCCTCTTGCT	3720
CAGCAGGCAC TTCTGAAAC AGGAAACAAG GAGAGTGACC TCCTAGCTTC ACTAGGACTA	3780
ACAGCTTTCT TCCTGGTCT GTTACGCTA GGGAAAAAGA GAGAACATA AGAGAAGAAT	3840
TCTAACACATT TGATTTGTA AAAATGGCTC TTTGTCAACT GTAGTGGTT GAAGTCAGCT	3900
AAGCTCGAGA AAGGACAAAT TTTGCTCTT CTTTTTGAT ATTCAAGAGCG ATAAAAATCC	3960
GTTTTTGAA GTTTCAAAG TTCCGAAAC CAAAGGCATT GCGCTTGATA AGTTTGATGA	4020
GATTATTGGT CGCTTCCAAT TTGGCGTTAG AATAGTGTAG TTGAAGGGCG TTGACGATTT	4080
TCTCTTGTC CTTTAGAAAG GTTTAAAGA CAGTCTGAAA AAGAGGATGA ACCTGCTTTA	4140
GATTGTCCTC AATGAGTCCG AAAAATTCT CCGGTTCCCTT ATTCTGAAAG TGAAACAGCA	4200
AGAGTTGATA GAGCTGATAG TGATGTTCA AGTCTTGTGA ATAGCTCAA AGCTTGTAA	4260
AAATCTCTTT ATGGTTAAA TGCATACGAA AAGTAGGGCG ATAAAAATGT TTATCGCTGA	4320
GTAAACG	4327

(2) INFORMATION FOR SEQ ID NO: 118:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 3521 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 118:

CTCTGGCCCT GCCACTCCAA CGTTTGTCA GGGTGTCTTT TTCATAAAGG AGTTCTTATG	60
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843

TTAGATATCA AACGTATTCG TACAGATTT GAAGCTGTCG CAGAAAAATT AGCTACACGT	120
GGTGTAGATG CTGCTGTCTT GAATGAAATG AAAGAAATCG ATGCTAAACG TCGTAACATC	180
TTGGTCAAGG TTGAAACTCT CAAAGCAGAA CGTAACACAG TTTCTGCTGA GATTGCCAA	240
GCTAAGCGCA ACAAGGAAA TACAGATGAC AAGATTGCTG CCATGCAAA TCTATCTGCT	300
GAGGTTAAAG CCTTGGATGC TGAATTGGCA GAAATCGATG CTAAATTGAC AGAATTACA	360
ACGACTCTTC CAAATATCCC AGCTGACAGC GTTCCCTGTTG GGGCTGACGA AGACGACAAT	420
GTGGAAGTTC GCCGTTGGGG TACTCCACGC GAGTTTGACT TCGAACCTAA AGCTCACTGG	480
GATCTCGGTG AAGACCTTGG TATCCTTGAC TGGGAACCGCG GTGGTAAAGGT AACAGGCCT	540
CGCTTCCCTCT TCTATAAAGG CCTCGGTGCT CGMTTGGAAC GTGCTATCTA CAACTTTATG	600
TTGGATGAAC ATGGAAAAGA AGGCTATACT GAAGTCATCA CACCTTACAT AGTCAACCAT	660
GATTCTATGT TTGGTACTGG TCAGTATCCA AAATTTAAGG AAGATACTTT TGAACTCAGC	720
GATACCAACT TTGCTTGAT TCCAAC TGCTGACT GAAGTTCCCTC TGACAAACTA CTACCGTGAT	780
GAAATCTTAG ACGGCAAAGA TCTTCCAATC TACTTCACTG CCATGAGTCC GTCATTCCGT	840
TCTGAGGCTG GTTCTGCCGG TCGTGATACG CGTGGCTTGA TCCGTTGCA CCAATTCCAC	900
AAGGTTGAAA TGGTCAAATT TGCCAAACCA GAAGAATCTT ACGAAGAATT GGAAAAATG	960
ACAGCCAACG CTGAAAACAT TCTTCAAAAAA CTCAACCTTC CATAACCGTGT CGTTGCTCTC	1020
TCTACTGGAG ATATGGCTT CTCAGCTGCG AAGACTTACG ACTTGGAAAGT GTGGATTCCA	1080
GCACAAAACA ATTACCGTGA AATCTCAAGC TGTTCAAACA CAGAAGATT CCAAGCCGT	1140
CGTCCCCAAA TCCGTTACCG TGATGAAGCA GATGGCAAGG TGAAACTCCT TCATACCTTG	1200
AACGGTTCTG GACTTGCAGT TGGACGTACA GTGGCTGCAA TTCTTGAAA TTACCAAAT	1260
GAAGATGGTT CTGTGACCAT CCCAGAAGCA CTTCGTCCAT ACATGGGTGG AGCTGAAGTC	1320
ATCAAACCAT AAAAATAAG GTTTAGCTAT TTCTAGCTAG ACCTTTTTTC GTAACCAAAT	1380
CAGATAAGCA CCTAGTACAA AGAATAAAAT AGTTAGGCAT ATAATGGTTT CAGCCAATAC	1440
CAGGTAATCC AGAAATGGAA GTTTCAAAT TCCCCTGAGCC ATCTTGAGCG AGGTCGCTGT	1500
GATAATGGTT GGGAAAGGTGA GGGCTGAGAA GGCTGGTTGA AAACCTTGTT TTAAAATGTT	1560
GGGCAGACGA GTTAAAACAA AGAAAAAGAA GGATTGAGAA GCCAAATCA TGACAATCAA	1620
GACCCAAGTC GGCAGGCTGG TTCCCTCTAC TCGAACTAGA GAAGCCAAGA GTAGAGAGAA	1680
AGGAGCACAG TAGATTCCTT CTTGTCCAAG CAAGGCTAGT GGGAGTGGAT GTTCTTTAA	1740
ATCGCTATAA ATAAGGGGAT AGAGATAGAA GGTCAAGAGA AAACCAAAAC TCAAGGTCGC	1800

844

ATAGGCATT TCGATAATAC CTACCAGAGG ATAGGTCAAG GCAGCCACTG	CTATCCCCAC	1860
ATAGAGAACC GTCCAGCTTG GAGTGGCATG AACCTCCGC CCTGGACAAG CAAACTTGAT		1920
GGTAAAACCA GCAATCAAGG TCAAATCCAA GAGAAATGAA AACACCAAA TCCCCTGTGC		1980
TACCAAAGGA AGATAAGAGA ATACCGAAA GACATAGGTC GATAAAATCA TCCCAGCCAT		2040
AGGAAAGGTT GCCATTCTG ACAAAAGAGG GGGCTTGGTC AATTCTTGCT TGGTTTCTT		2100
CCAATTAAAG AGATGCAGAA TTAGAAAGTA ATACCATAAA ACCAAACCAA TCAGACTAA		2160
AAGATGGAT AGAACCGGCA ACCTATCTAA ATAAGGATTT CCAGCTCCTG CCAAACCTAG		2220
CAAACAACT GAAAATACTA AGGGGAGTTT TTTCATCCTA ACCTCCAATA ATCATGTTAG		2280
TTTCAGTATA ACATAAAAGC GCTTAAATGA GGATTTAAAA AAACGAGTCC GCTTATTTCA		2340
GACTTCATTT TACTCAGATA TGAATTAGGC ATAAGGTTGC AATTCTGGAT TAATTGGTGT		2400
ATTAGCTAAG TTGTTGGCAT AGTTACAGAG GATTGCTAGG CTGACACCAA AAACACACATC		2460
CAAGGCATTT TGTTGAGTGT AGCCAGCTTC TAAAAACTCA GACAAGGCTT CATCTCCTAC		2520
ACGACCCCTG GTATTGATAA CTGCCAAGGT AAACCTAGCT AGGGTATCCA ATTTAGGATC		2580
TGTTTCATT GGAGTACGAT TGCGAAGAGC TTGAATCAAG TCATCATTCA TCTGGATTTG		2640
TTTGATGGAA AAGGCTGTGT GACCTGCGAC ACAGAAGGCA CAACCATTGG TCACGGCTGC		2700
CGTGATTTGC ACCACTTCAC GCTCAACGGG TGTCAGGCTG TTGCGACGGT GGATAGATGA		2760
GACAATTGG TAGGCTTCTA AAACAGTCGG GGCATTGGCC AAGAGACCGA TTAGGTTGGG		2820
AATATAGCCA TTGTTGTCTT TTTCTACTGT TTCAAGAATT TCTTTCACTT CTGCTGGTGC		2880
TGACTCTACT GTATGGATAG TAAATGTTGT CATAAGATAC CTCTTTCTT ATTATTGACA		2940
CTAATATTAT TGGAAAATCT TATAAAATCC TGATTCCCTAA GTTTATCTAA GATAAGCTT		3000
TATTCTCTCA TAAGATTTTC GTGTTATAT TAGTTTATCA CACTTCCAAT CACTTGTATA		3060
ATATATATTA TATATCGGC TGATAAAAAT TATTTATAGG CAAAAAAATC ACACGAGCTG		3120
TGTGATTCCA TTATTTGTCA AAATACTTTT TAGTTTCAGC AATAACGACT GGCACAAAGA		3180
CCAAGAGGGC AATCAAGTTT GGAGAGGCC TCAAGGCGTT AACGATATCT GCGATAATCC		3240
AGACCATATC CAACTCGATA AATCCTCCTA ACAAGACCAT GAGCACACAAA ACCACACGGT		3300
AGAGCCAGAT AAAGCGAACCC CAAAGAGGA ACTCAAAACA GCGTTCTCCG TAATAGTTCC		3360
AACCTAGAAT CGTTGAAAG GCAAAAGTA CAAGGAAGAT GGTCAAGAGA GCAGGCCCAA		3420
AGTGTGAAAA GTTTGTTGAG AAAGCTGACT GAGTCAGGC AACCCCATTC AAGTCACCGC		3480
TCCAAACTCC AGTTACCAAG ATGGTCAAAC CAGTTAGAGT A		3521

(2) INFORMATION FOR SEQ ID NO: 119:

845

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 1968 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 119:

AACCTGGCA AGCAAGCTAA AAGCAATGGG ACCTGGAATC CTAATGGCAA CTGCCGCTGT	60
TGGAGGTTCC CACATTGTAT CCTCAACTCA AGCTGGCGGT TCTTACGGTT GGTCCTACT	120
TCTCTTGTC ATCTTAGCCA ATGTCCTTAA ATATCCATTT TTCCGTTTTG GTGCTGAATA	180
CACAGCTGAT ACTGGAAAGA CTTTGGTTGA AGGTATGCC GAAAAAGGAA AACTCTATCT	240
CTGGATTTTC TTTATCCTCA ATGTCCTTTC GGCTATGGTC AACACGGCTG GTGTTGCCAT	300
TCTGTGCTCA GCTATCATCG CCAGTGCCCTT CCCAATGATT GGACTTAGCA TTACTCAGTG	360
GTCCTCATT CTCGTTGCAA TCATTTGGGC TATGCTACTC TTTGGAGGCT ACAAACTTTT	420
AGACGGCATG GTCAAATGGA TTATGTCTGC CTTAACCAATT GCGACTGTTT TTGCAGTTT	480
CATTGCGGCG GTCAAGCATC CAGAAATACAG TTCTGATTTT GTCGAGAAGA CACCTTGGCA	540
AATGGCAGCT CTGCCCTTC TCGTCTCCCT CCTAGGATGG ATGCCGGCTC CTATTGAAAT	600
TTCAGCCATC AATTCACCTT GGTCAAGCTGA AAAGAGAAAG ACCGTCAACT TTAACACAGA	660
AGACGGCTTG TTGACTTTA ACACGGTTA TATTGGAACA GCTATCCTAG CCGCTTCTT	720
TGTGGCACTG GGAGCACTGA TTCAGTATCC TACAGGGCAG GCGGTTGAAG CTGCTTCAGC	780
CAAATACATC TCTCAATTGCA TGGGCATGTA TGCCCTCTGTT CTTGGCGAAT GGTCCCCTTA	840
CTTGATTACC TTTATTGCT TCCCTCTGTAT CTTTGGAAACA GTTATAACTG TTATCGATGG	900
CTATTCTCGC GTTAATCAGG AATCTCTCCG ACTGCTAATC AGTCAAAAG AGGACAATCG	960
TAAATCTTG AACATCTGGA TGACCATCAC TGCTATCATC GGTATCGTCA TTATCAAGTT	1020
CTTCGCTGGT CAGGTTTCAA CCATGCTCCG CTTTGCCATG ATTGGCTCTT TCCTGACAAC	1080
ACCTTTCTTT GCTCTTTGA ATTACGGCTT GGTAAACGGGT GAAAACAAAA ATCTTCTTC	1140
TTGGCTCAAA CACCTTGCCA TTGCGGGATT GATTTCTCTC TTTGCTTCGC CATCTTCTTT	1200
ATCTACGCAC TCGCAATCGG AAAAGCAGGG TAAGGGACAA GCGCGAGATG AAGATAAGGT	1260
TTCATTTCAA GAGAAAATTC AGCAAATATT TCTATGATAA AAAGCATAAG ACAAGGTTT	1320
TGAAGACCTG AACTPATGCT TTTTTACGTT CTTAAAGACT GTTTATACTC AAAAAACAGT	1380
TGAACAACTT CAACCCACCTC TTATAAGAAC TTTTAACTAT TCGAGAATCT CTTCAAAACCA	1440

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CGTCAGCTCT ATCTGCAACC TCAAAGCTGT GCTTTGAGCA ACCTGCGACT AGCTTCCTAG	1500
TTTGCTCTTT GATTTTCATT GAGTATTAAT TCTCCTTTTC CAAACTCATAC AAATCTGCCA	1560
TAATAGCTGC GACATGTTG ATATCTTCCA GCATGCCCTCG CATTTCAAAG TCAGCCAATA	1620
CAGGGAAGCC AAAGCGTTGA CTGTATTGCT TGGCTGTTAG GCAGTATTGG TTATTAAGT	1680
TACGATTCC TGACCCAACC ACACCAAAAC ACTTACTAGC ATTGTTACCA TAGGCAATAA	1740
AATCTCCCAC CGGTGTCGTC AAAATCTCAA CATCTCCGTT ATCCACGCCA TTCCCACCTT	1800
CGAGATAGGT CGGCAAAAAA GCGACATAGG GATGGTCCAT TTCATAGAAA TTTTGCCCTT	1860
CCTTGACCAA ATCCCTGATA TGAATCTTT GAACCTCAAT CCCTTGTAC TGGGACAAGA	1920
GATAGTCTTT CAAGCGCGTC ACAAAACTTT CAGTGTGCC ACTCAAGG	1968

(2) INFORMATION FOR SEQ ID NO: 120:

- (i) SEQUENCE CHARACTERISTICS:
- (A) LENGTH: 7172 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 120:

CCGCATTTTT TATCACTAGA CTCGAGACAT CTTTGAGTG GCTCTTGCTC TCTGGTTAA	60
TTTCTTCCT TGCTCAAGGA CTCCCTGCTAT TTCTCTTGGT CGTCCGACTC AAACATCAAT	120
TCGCTGAGAT TTATCCTCAA ATCAAAAAA AGATTCGCTT CTACTATTAA GGGGTTCTCA	180
CCATTGATTT TCTATTTTTT GTTCTCTTAG CCTTCATTAG TTCTCAGCGT TTTTCATCTC	240
TTATGCCAAT CATCACTGCT TGCCATTCTA CTTTTTATTAA TATGACAGCT GACTACCTAA	300
GAGAAAACTA TCCAGACTTT TACGACAAAC ACATCTCTT ATGGGAGTGT CTCTAAAGAA	360
AAGGGGGTTT TAGCATGAAA AAAATCATCT TCATCAAAAC CATTCAACTC CTTGTCATTG	420
ATGGAATCAT GCTGGCATTT TTGACATTAA AAAGGGGGCT TACTTGGAC TGGATTTGA	480
TTTATAGCGG TTGGCTCAT TTCTTTCATC CTGTGCTATT GACCTATCTT TCAAACCAAC	540
TTTGTGACCA CTTTAGTTAA CTCTATTCCC AGATTAGACC GAGATTCTGG CGTTTGCTT	600
TACAAATTCT CCTATGGGAT AGCCTGATGA TTCTCTCCCTT GGTGTCTTAA AGTGTATTC	660
CACTTTCCCT TCAGGAACT CTCCCTCATCC TAGGACATCT CATCCCTTCC TATCGCATCT	720
GCCAAAGCCT GAAAAGAGAC TTCCCCAAG CATATCAAGA ACCGATTTCT TTTTGGAGTA	780
TTTTATGATA GATGAGAAAG ACCAAGCCGA CTGGGCTTGG TCTTTCTTAT CTCTTTTAG	840
TATCTAGGAT AATGGTAACA GGTCCATTAT TAACCAGCTC AACCTGCATA TCTGCTCCAA	900

847

AGATGCCCTGT	CTGAACGGGC	ACTTCTTGCG	CTAATTTTG	ATTGAAAGCA	TCATAGAACGT	960
CTGATGCCAT	ATCAGGTTA	GCTGCCCTG	TAAAGgCTGG	ACGATTGCCT	CTCTTAGTAT	1020
CCGCAAAGAG	GGTAAACTGA	GAAATAGAGA	GGATTTCTCC	TTCAATATCT	TTGACAGACA	1080
GGTTCATCTT	GCCTTCTGCG	TCTGAAAAAA	TCCGCATATT	GACCAGTTT	CTCACAGCAT	1140
AGTCCAATC	TTCCTCTTGG	TCCTCTGGTC	CAACACCAAC	CAGCAATAAA	AGTCCCTGAT	1200
TGATTTTCC	CTGAATCTGG	CCTTCTATAC	TCACTTGGGC	TTTTTTAACCC	CGTTGGATAA	1260
TGATTTTCA	AATAGCCTT	CTAGTAAGAG	CTAGGACAAC	TAGCCGTTGG	TCCGTTTGAC	1320
AGAGTAAACT	TCTGGCACAC	TCTTAATTTC	ATCGACAACC	GTGGTCAGTG	TAGAGAGGTT	1380
GGCAATACCG	AAGgACACAT	GGATATTAGC	AAACTTCATA	TCCTTGGTTG	GTTGGGCATT	1440
GACCGTTGAA	ATATTCTTGG	TTGTATTTGA	AAGAACTTGC	AGTACATCGT	TCAACAGTCC	1500
TGTACGGTTG	AGACCGTAGA	TATCGATATG	GGCCATATAC	TCCTTATTTC	AGCTAGGGTA	1560
CTGGCTTCC	CATTCCACAT	CAAGGAGACG	TTGCTCGTAG	TTTTCTTGGG	CACGCAGGTT	1620
CATACAGTCC	ACACGGTGAA	TAGCCACACC	ACGACCCCTG	GTAATGTAGC	CAACAATATC	1680
GTCACCAGGC	ACGGGGTTAC	AACACTTAGC	AATCCGCACT	AGGAGACCAAG	AAGCACCTTC	1740
AATAACCACT	CCCCCTCAT	GCTTGACCTT	GAGGGTTCT	TTATTTTCAA	CCTTGACCTC	1800
GCCACCTTTC	ACAAGCTCCT	CTGCCCTCAGC	TTTGGCCTTG	GCACGGCTCTT	CCTCACGGCG	1860
TTCCTTTCA	GTCAGACGGT	AAAGACGGT	AATCGCACCG	ATTTCCCCAA	AACCAATGGC	1920
CGCAAAGAGG	GAGTCTTCTG	TCTTGTAACT	GGTCTTTGC	AGAACTTGAT	CCATGTGGCG	1980
CTTGTCCATA	AATTTATTTG	CCACATAGCC	ATTTCTTGG	AACTGAGCCA	TCAGCATCTC	2040
ACGACCCCTTG	TTGACAGACA	ATTCCTTATC	TTGGTTTTA	AAGAACTGGC	GAATCTTATT	2100
GCGCGCCTTG	CTAGTCTTGA	CCATATTGAG	CCAGTCACGG	CTAGGTCCAA	AGGAGTTGG	2160
GTTGGCGATA	ATTTCAACCT	GATCCCTGT	CTTTAACCTG	GTTGTCAGTG	GAACCATGGC	2220
GCCATTGACC	TTGGCACCAAG	TTGCTTTTC	ACCGACCTTG	GTATGGATT	CGTAGGCAA	2280
ATCAATCGGT	CCTGAATCTT	TGGAAGGGA	ACGGACAGCT	CCATCTGGGG	AAAAAACGTA	2340
AATCTCCTCA	GCCAAATAGT	TTTCCTTAAC	AGAGTCCACA	AATTCTTAG	CATCATCAGC	2400
CTGGCTTGG	AGCTCCATCA	TCTCCTTGAT	CCAGTTCATT	CCAATAGCTG	ATTCCCTTGCT	2460
GTAACTTGC	CCCTTATAC	CTTCTTATA	AGCCCAGTGA	GCCGCAACCC	CGTACTCAGC	2520
CACCTCGTGC	ATTTCTTGG	TTCGAATCTG	GAATTCAATC	GGCCCTTTG	GTCCATAAAC	2580
AGTCGTATGG	ATAGACTGAT	AACCATTGGC	CTTGCAGTTG	GCGATATAGT	CTTTGAAGCG	2640

848	
ACCTGGCATC GGTTCCAAA ATTCATGCAC GTAACCAAGC ATGGCATAAA CATCACTTGT	2700
GGTATCTAAA ATACAACGAA TAGCAATCG ATCATAGATT TCCTCAAACC GTTTTCTCTT	2760
GTCCTGCATT TTGCGGAAAA TTGAGTAAAT ATGCTTGGGA CGACCATAAA TCTTCCCTTT	2820
CAAGTGACGT TCTGTCGTAT ACTCCTCTAA TTTTGTGACT ACCTCATCCA CCAAGGCCCTC	2880
ACGCTCCCTG CGCTTTCCCT TCATCATATG GGTAATCTTG TAAAACCTCCG TTGGATTGAG	2940
ATAACGGAAA GACAAGTCTT CTAATTCCCA TTGACACTG GAAATCCCCA AACGATGGGC	3000
AAGCGGGCA TAGATTTCCA TGTTTCTTT GGAAATACGC TCCTGCTGT CTTTTCGAAG	3060
ATGTTTCAGG GTCGCATAT TGTGCAAGCG GTCAAGACAGT TTGACCAAAA TAACGCGGAT	3120
GTCCTCAGAC ATGGCCATGA GCATCTTGCATG ATGATTTCC GCTAATTGCT CCTCGATCGA	3180
TTTGTACTCG ACCTTGCAAA GCTTGGTAAC TCCGTCAACA ATCATCCGCA CATCAGGACC	3240
AAACTCTCTT TCCAATCGT CCAAAGTCGC ATCTGTATCT TCCACCACAT CATGCAAGAA	3300
TCCACAAGCT ACTGTTACAG CATCCAGCTT TAGCTTAGCT AAAATACCTG CCACTTGGAT	3360
AGGGTGAATG ATATAAGGCT CGCCTGATTT GCGATATTGA CCACGTGGC ATTCAACAGC	3420
ATAGACCAAG GCCTTATGGA CAAAATGAAC ATCCTCTTCC GTTAAATATT CTTTGGTTAA	3480
AGCGACAAC TCTTCGCTG TTAAATTACAC TTCTTTCGGC ATCTCTACTC TCCAATTCTT	3540
CCTACCATT TATCACTTTT TTAAGAATAT GAAAACCTAGA TTGGAACAGA ATAAGAAAAAA	3600
AATAATTCAA AATTGCTTGA TAATTCTGAA TTATTGGTCC GTAATATACT ACCAAGTTAG	3660
ATTTTAAACT TAGGTGATAG AAGGAGAGAT AGAAGAACCG AAACCATATT GTAACCCAAA	3720
GACTTTCTGA CTTCCCCAAT TCCATTGAAAG ATACGAAAGA TAAACGGTGG AACTCGTATC	3780
ACATACACTG GTACCTTGAC TGGATTTGG ATTAATACT AAATGAAAAT CAAAGAGCAA	3840
ACTAGGAAAC TAGCCGCAGG TTACTCAAAG CACCGCTTGT AGGTTGCAGA TAAAGTTGAC	3900
CGGGTTGAA GAGATTTTG AAGAGTATAA AAATCCTCAA GATACTTTCT TCTATCCTTT	3960
AGTTTATAAG GAGAATACCT ATGAAAAAAA CTGCTATTTC TATCTTGTCT CTCCTAATGT	4020
TAGGAGTTG CTGCCTGTTCTTATTCAGCC AGCAAAGCTA TAAAAAACAG TCGTTCAATA	4080
CTATGCTAAC GACCAGAACCC TGCCCAGTAG GATAACTTAT AGTGAATATA GCGACAAATG	4140
AGAAGCCAAAC TACGGTAGCA CTCTAAACAT CACGTCTATC AAACAAGCTA ATGACGGAGT	4200
TTATGCAACC TATGAAGGGC AATTGACACC TTTCCAATAT TGATAAATTG ATAACCAGCC	4260
TGTCTTCATC TAGTCATGCT GGTTTTAAG TTCATTAAAC ATCCTTACCT ATTCTCCCTA	4320
ACTGTGCTAT ACTTAATTAA TACTCAATGA AAATCAAAGA GCAAACCTAGA AAGCTAGCCG	4380
CAGGCTGTTCAAAAGCAGCTGCTTCAAGT TGACGCGGTT TGAAGAGATT	4440